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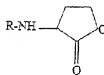
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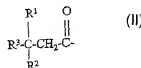
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(54) Title: MODULATION OF STAT ACTIVITY



(I)



(II)

(57) Abstract: The use of a compound selected from a range of compounds including quorum sensing molecules, N-acyl homoserine lactones, N (3 oxododecanoyl) L homoserine lactone, inhibitors to modulate STAT activity for the treatment of a range of diseases including cancer, breast cancer, obesity, lipid metabolism disorders, immune disease, immune deficiency or immune disorders. The range of compounds also include compounds of formula (I) in which R is an acyl group of formula (II).

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MODULATION OF STAT ACTIVITY

Field of the Invention

The invention relates to the modulation of STAT activity. The invention also relates to compounds preferably (though not exclusively) quorum sensing molecules such as those produced by *Pseudomonas aeruginosa* for inhibiting STAT activation.

Review of the Art known to the Applicant(s)

STATS (Signal Transducers and Activators of Transcription) are evolutionarily conserved molecules (proteins) identified in the genomes of mammals, flies, worms and even the slime mould *Dictostelium*. Inactive STAT proteins are cytoplasmic or associated with membrane

growth factor and cytokine receptors. Ligand binding to these receptors causes the tyrosine phosphorylation of associated STATs, their homo- or heterodimerisation and translocation to the cell nucleus, where they interact with promotor elements to activate target gene expression.

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Biological functions of mammalian STAT proteins have been revealed by gene targeting experiments in mice. STAT1 knockout mice show defective macrophage function and sensitivity to viral infection, while the absence of STAT5a and STAT5b causes defects in T cell growth. Notably, mice lacking STAT4 are defective in Th1 responses and STAT6-
10 deficient mice are defective in Th2 responses. Deletion of the gene for STAT3 results in early embryonal lethality. This is possibly due to the singular role of STAT3 in the proliferation of several cell types. Conditional deletions of STAT3, however, demonstrate a requirement in T cells for IL-2 and IL-6-induced proliferation, in macrophages to counteract chronic inflammation and in keratinocytes for wound healing. Thus, consistent with findings that link
15 the invertebrate STAT protein to immune function, a common theme in STAT knock-out mice is the disruption of aspects of immune function.

Summary of the Invention

20 In its broad concept the invention provides an autocrine/paracrine signalling pathway which is associated with cell growth/proliferation, modifications to which can be used to alter cell growth and/or cell proliferation.

In one aspect the invention provides an autocrine/paracrine signalling pathway which
25 activates STAT wherein the pathway requires JAK activity and does not require Erb1 activity and is not induced by EGF.

In a further aspect the invention provides a process wherein STAT dimmers accumulate in the cytoplasm wherein the process does not require Erb1 activity or JAK activity.

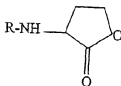
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In a further aspect the invention enables the modulation of any one of these pathways or processes to alter the amount of activated STAT.

In yet a further aspect the invention encompasses the use of a compound selected from:-

- 5 JAK, ErbB1, EGF, ErbB1 inhibitors, EGF inhibitors, STAT inhibitors, interleukin- 13 (IL-13), IL-13E13K (IL-13 in which the Glu at position 13 is substituted by a Lys residue), sulphur methoxazol, ubiquitin E3 ligase, serine phosphatase, tyrosine phosphatase, SOCs, Pias proteins (protein inhibitors of activated STAT), STAT1 inhibitors, STAT2 inhibitors, STAT3 inhibitors, STAT4 inhibitors, STAT5A inhibitors, STAT5B inhibitors, STAT6
- 10 inhibitors, JAK inhibitors, AG 490, α -amanitin, transcription inhibitors, quorum sensing molecules, N-acyl homoserine lactones, N-(3-oxododecanoyl)-L-homoserine lactone, oxygen radical scavengers, N-acetyl Cysteine (NAC), diphenylene iodonium chloride (DPI), inhibitors of COX1, inhibitors of COX2, aspirin, ketorolac, indomethacin, or panCOX inhibitors to modulate STAT activity for the treatment of cancer, breast cancer, multiple
- 15 myeloma, head and neck cancers, leukaemia, HTLV-1-dependent leukemia, large granular lymphocyte (LGL) leukaemia, erythroleukemia, acute lymphocytic leukemia (ALL), chronic lymphocytic leukaemia (CLL), acute myelogenous leukemia (AML), chronic myelogenous leukaemia (CML), megakaryotic leukaemia, lung cancer, renal cell carcinoma, prostate carcinoma, melanoma, ovarian carcinoma, pancreatic adenocarcinoma, lymphoma, EBV-
- 20 related lymphoma, Burkitt's lymphoma mycosis fungoides lymphoma, HSV saimiri-dependent (T cell) lymphoma, cutaneous T cell lymphoma, obesity, lipid metabolism disorders, immune disease, immune deficiency or immune disorders.

In another aspect the invention relates to the use of a compound of the formula I



in which R is an acyl group of the formula II



wherein one of R^1 and R^2 is H and the other is selected from OR^4 , SR^4 and NHR^4 , wherein R^4 is H or 1-6C alkyl, or R^1 and R^2 together with the carbon atom to which they are joined form a keto group and R^3 is a straight or branched chain saturated or unsaturated aliphatic hydrocarbonyl group containing from 8 to 11 carbon atoms and is optionally substituted by one or more substituent groups selected from halo, 1-6C alkoxy, carboxy, 1-6C alkoxycarbonyl, carbamoyl optionally mono- or disubstituted at the N atom by 1-6C alkyl and NR^5R^6 wherein each of the R^5 and R^6 is selected from H and 1-6C alkyl or R^5 and R^6 together with the N atom from a morpholino or piperazino group or any enantiomer thereof with the proviso that R is not a 3-oxododecanoyl group to modulate STAT activity for the treatment of cancer, breast cancer, multiple myeloma, head and neck cancers, leukaemia, HTLV-1-dependent leukemia, large granular lymphocyte (LGL) leukaemia, erythroleukemia, acute lymphocytic leukemia (ALL), chronic lymphocytic leukaemia (CLL), acute myelogenous leukemia (AML), chronic myelogenous leukaemia (CML), megakaryotic leukaemia, lung cancer, renal cell carcinoma, prostate carcinoma, melanoma, ovarian carcinoma, pancreatic adenocarcinoma, lymphoma, EBV-related lymphoma, Burkitt's lymphoma mycosis fungoides lymphoma, HSV saimiri-dependent (T cell) lymphoma, cutaneous T cell lymphoma, obesity, lipid metabolism disorders, immune disease, immune deficiency or immune disorders.

In a further aspect the invention relates to any one of these uses wherein the R group is selected from

5



and

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10 wherein R³ is as defined above.

In another aspect the invention relates to any of these uses wherein the group R³ is an 8-11C straight or branched chain alkyl group optionally substituted by a substituent selected from bromo, carboxy and methoxycarbonyl.

15

In a further aspect the invention relates to any one of these uses wherein the R³ group is such that the group R in formula I is selected from;

- 3-oxoundecanoyl;
- 11-bromo-3-oxoundecanoyl;
- 20 10-methyl-3-oxoundecanoyl;
- 6-methyl-3-oxoundecanoyl;
- 3-hydroxydodecanoyl;
- 12-bromo-3-oxododecanoyl;
- 3-oxotridecanoyl;
- 25 13-bromo-3-oxotridecanoyl;
- 3-hydroxytetradecanoyl;
- 3-oxotetradecanoyl;
- 14-bromo-3-oxotetradecanoyl; and
- 13-methoxycarbonyl-3-oxotridecanoyl.

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In a further aspect the invention relates to any one of these uses wherein the R^3 is an 8-11 straight or branched chain alkenyl group optionally substituted by a substituent selected from bromo, carboxy and methoxycarbonyl.

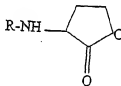
- 5 In a further aspect the invention relates to any one of these uses wherein the R^3 group is such that the group R in formula I is selected from;

3-oxo-12-tridecenoyl;
3-oxo-7-tridecenoyl;
3-hydroxy-7-tetradecenoyl;
10 3-oxo-9-tetradecenoyl;
3-hydroxy-9-tetradecenoyl;
3-oxo-10-tetradecenoyl;
3-hydroxy-10-tetradecenoyl;
3-oxo-11-tetradecenoyl;
15 3-hydroxy-11-tetradecenoyl;
3-oxo-13-tetradecenoyl; and
3-hydroxy-13-tetradecenoyl.

In another aspect the invention relates to the use of JAK, ErbB1, EGF, ErbB1 inhibitors, EGF
20 inhibitors, STAT inhibitors, interleukin-13 (IL-13), IL-13E13K (IL-13 in which the Glu at position 13 is substituted by a Lys residue), sulphur methoxazol, ubiquitin E3 ligase, serine phosphatase, tyrosine phosphatase, SOCs, Pias proteins (protein inhibitors of activated STAT), STAT1 inhibitors, STAT2 inhibitors, STAT3 inhibitors, STAT4 inhibitors, STAT5A inhibitors, STAT5B inhibitors, STAT6 inhibitors, JAK inhibitors, AG 490, α -amanitin,
25 transcription inhibitors, quorum sensing molecules, N-acyl homoserine lactones, N-(3-oxododecanoyl)-L-homoserine lactone, oxygen radical scavengers, N-acetyl Cysteine (NAC), diphenylene iodonium chloride (DPI), inhibitors of COX1, inhibitors of COX2, aspirin, ketorolac, indomethacin, or panCOX inhibitors, for the preparation of a medicament for the treatment of cancer, breast cancer, multiple myeloma, head and neck cancers, leukaemia,
30 HTLV-1-dependent leukemia, large granular lymphocyte (LGL) leukaemia, erythroleukemia,

acute lymphocytic leukemia (ALL), chronic lymphocytic leukaemia (CLL), acute myelogenous leukemia (AML), chronic myelogenous leukaemia (CML), megakaryotic leukaemia, lung cancer, renal cell carcinoma, prostate carcinoma, melanoma, ovarian carcinoma, pancreatic adenocarcinoma, lymphoma, EBV-related lymphoma, Burkitt's lymphoma mycosis fungoides lymphoma, HSV saimiri-dependent (T cell) lymphoma, cutaneous T cell lymphoma, obesity, lipid metabolism disorders, immune disease, immune deficiency or immune disorders.

In a further aspect the invention relates to the use of a compound of the formula I



in which R is an acyl group of the formula II



wherein one of R^1 and R^2 is H and the other is selected from OR^4 , SR^4 and NHR^4 wherein R^4 is H or 1-6C alkyl, or R^1 and R^2 together with the carbon atom to which they are joined form a keto group and R^3 is a straight or branched chain saturated or unsaturated aliphatic hydrocarbonyl group containing from 8 to 11 carbon atoms and is optionally substituted by one or more substituent groups selected from halo, 1-6C alkoxy, carboxy, 1-6C alkoxy carbonyl, carbamoyl optionally mono- or disubstituted at the N atom by 1-6C alkyl and NR^5R^6 wherein each of the R^5 and R^6 is selected from H and 1-6C alkyl or R^5 and R^6 together with the N

atom from a morpholino or piperazino group or any enantiomer thereof with the proviso that R is not a 3-oxododecanoyl group for the preparation of a medicament for the treatment of treatment of cancer, breast cancer, multiple myeloma, head and neck cancers, leukaemia, HTLV-1-dependent leukemia, large granular lymphocyte (LGL) leukemia, erythroleukemia, acute lymphocytic leukemia (ALL), chronic lymphocytic leukemia (CLL), acute myelogenous leukemia (AML), chronic myelogenous leukemia (CML), megakaryotic leukemia, lung cancer, renal cell carcinoma, prostate carcinoma, melanoma, ovarian carcinoma, pancreatic adenocarcinoma, lymphoma, EBV-related lymphoma, Burkitt's lymphoma mycosis fungoides lymphoma, HSV saimiri-dependent (Tcell) lymphoma, cutaneous T cell lymphoma, obesity, lipid metabolism disorders, immune disease, immune deficiency or immune disorders.

Description of the Drawings

The invention will be described by way of example with reference to the accompanying drawings in which:

Figure 1 is a diagrammatic representation of the autocrine/paracrine pathway which activates STAT3.

Figure 2 shows:

ErbB and STAT protein expression in BC cell lines. Lysates were prepared from BT20 (lane 1), MCF-7 (lane 2), T47D (lane 3), MDA-MB-231 (lane 4), MDA-MB-468 (lane 5) and BR293 (lane 6) cells. For ErbB1, STAT1 and STAT3 200 µg protein from each cell lysate were used for Erb2 and Erb3 400 µg of protein was used and separated by SDS-PAGE, transferred to PVDF membranes and probed with anti-ErbB (upper panel) or anti-STAT (lower panel) antibodies as indicated. One set of lysates was used throughout.

Figure 3 shows: Tyrosine phosphorylation of ErbB proteins in BC cell lines. Lysates were prepared (see Materials and Methods) from BT20 (lanes 1, 2), MCF-7 (lanes 3, 4), T47D

(lanes 5, 6) MDA-MB-231 (lanes 7, 8), MDA-MB-468 (lanes 9, 10) and BR293 (lanes 11, 12) cells that had been serum-starved (-) or starved and treated with EGF (5nM) for 15 min (+). ErbB proteins were collected as immune complexes, separated by SDS-PAGE, transferred to PVDF membrane and probed first with an anti-phosphotyrosine antibody and subsequently with the corresponding anti-ErbB antibody, as indicated. ND indicates that the protein is not expressed at detectable levels by the cell line (see Figure 2). Numbers below each panel show the level of tyrosine phosphorylation, quantified with Image Quant software (Fuji) and expressed as the ratio $\alpha\text{PY} / \alpha\text{ErbB}$, whereby the unstimulated value for each protein in each cell line is set as 1. The results shown are compiled from several experiments in which ErbB proteins from each cell line were analysed at least three times with similar results.

Figure 4 shows:

Tyrosine phosphorylation of STAT1 and STAT3 proteins in BC cell lines. Lysates were prepared from BT20 (lanes 1 and 2), MCF-7 (lanes 3 and 4) T47D (lanes 5 and 6), MDA-MB-231 (lanes 7 and 8), MDA-MB-468 (lanes 9 and 10) and BR293 cells (lanes 11 and 12) that had been serum-starved (-) or starved and treated with EGF (5nM) for 15 min (+). Proteins (200 μg) were separated by SDS-PAGE, transferred to PVDF membrane and probed first with an anti-phosph-STAT1 or anti-phospho-STAT3 antibody and subsequently with the corresponding anti-STAT antibody as indicated.

Figure 5 shows:

EGF-induced formation of DNA complexes by STAT proteins in BC cells. Extracts were prepared from MDA-MB-468 (lanes 1-6), BT20 (lanes 7-12) and BR293 cells (lanes 13-18) that had been serum-starved (-) or starved and treated with EGF (5nM) for 15 min (+). Equal amounts (25 μg protein) of each extract were incubated alone (lanes 1, 4, 7, 10, 13 and 16) or with antibodies specific for STAT1 (lanes 2, 5, 8, 11, 14 and 17) or STAT3 (lanes 6, 9, 12, 15 and 18) and aradiolabelled oligonucleotide duplex corresponding to the M67 sequence derived from the *c-fos* SIE. STAT1 homodimers (1:1), STAT3 homodimers (3:3),

heterogeneous STAT3 complexes (3H) and supershifted STAT3 complexes (3SS) are indicated. In subsequent figures only the upper parts of the EMSA gels are shown.

Figure 6 shows:

- 5 Inhibition of EGF-induced phosphorylation of ErbB1 and DNA binding of STAT proteins.
(a) Serum-starved MDA-MB-468 cells were pre-treated with PD153035 (100nM) for the times indicated and then treated with EGF (5nM) for 15 min (+). Lysates were prepared, from which ErbB1 proteins were collected as immune complexes, separated by SDS-PAGE, transferred to PVDF membrane and probed with an anti-phosphotyrosine antibody as indicated. (b) Equal amounts (25 µg protein) of each lysate were incubated alone (lanes 1, 4, 7, 10, 13, 16 and 19) or with antibodies specific for STAT1 (lanes 2, 5, 8, 11, 14, 17 and 20) or STAT3 (lanes 3, 6, 9, 12, 15, 18 and 21) and a radiolabelled oligonucleotideduplex corresponding to the M67 sequence derived from the *c-fos* SIE. In this and subsequent figures only the upper parts of the EMSA gels are shown. (c) BT20 (lanes 1-6) and MDA-MB-468
15 cells (lanes 7-9) were serum-starved (-), treated with EGF (5nM) for 15 min (+) or pre-treated with AG490 (100 µM) for 30 min and then treated with EGF (5nM) for 15 min (+). DNA binding by STAT proteins was analysed as described in the legend to Figure 5. The STAT1 homodimer is indicated (1:1). (d) BT20 (upper), MDA-MB-231 (middle) and BR293 cells (lower panel) were incubated in serum-free medium for 24 (lanes 1, 2), 48 (lanes 3, 4) or 72
20 hours (lanes 5, 6). Cell extracts were prepared and equal amounts of each were incubated directly with the M67 DNA probe (lanes 1, 3, 5) or after pre-incubation with an anti-STAT3 antibody (lanes 2, 4, 6). STAT1 homodimers (1:1) and supershifted STAT3 complexes (3SS) are indicated.

- 25 Figure 7 shows:

Delayed activation of STAT3 involves autocrine/paracrine signalling.

- (a) Extracts were prepared from serum-starved BR293 cells (lanes 1 and 6) or starved cells stimulated directly with 10% FCS for the times indicated (lanes 2-5), or after pre-treatment with AG490. STAT3 and JAK2 proteins were collected as immune complexes, separated by

SDS-PAGE, transferred to PVDF membrane and probed first with an anti-phosphotyrosine antibody and subsequently with

anti-STAT3 or anti-JAK2 antibodies as indicated. (b) Serum-starved BR293 cells were stimulated with 10% FCS for the times indicated and activation of STAT DNA-binding was assayed with the M67 DNA probe in nuclear (left panel) and whole cell extracts (right panel). (c) BR 293 cells were serum-starved (lanes 1 and 2), or serum-starved and treated with 10% FCS for 2 hours (lanes 3 and 4).

Alternatively, after 2 hours in 10% FCS, cells were washed and incubated in serum-free medium for a further 4 hours, whereupon conditioned medium from the cells was transferred to fresh serum-starved BR293 cells, which were harvested after 15 minutes (lanes 5 and 6). STAT DNA-binding activity in nuclear extracts was assayed with the M67 DNA probe. (d) Extracts were prepared from serum-starved BR293 cells (lane 1) and cells stimulated with EGF (lane 2) or conditioned medium (lane 3) for 15 min. STAT3 proteins were collected as immune complexes, separated by SDS-PAGE, transferred to PVDF membrane and probed first with an anti-phosphotyrosine antibody and subsequently with an anti-STAT3 antibody as indicated.

Figure 8 shows:

Inhibition of serum-induced STAT DNA binding. (a) BR293 cells were left untreated (lanes 1, 2) or treated with 10% FCS for 2 hours in the absence (lanes 3, 4) or presence of 100 μ M AG490 (lanes 5, 6). In addition, conditioned medium from serum-starved cells (lanes 7, 8) or from cells incubated in 10% FCS for 2h (lanes 9-12) was transferred for 15 minutes to fresh serum-starved cells (lanes 13-16) or starved cells pre-treated with 100 μ M AG490 (lanes 17, 18). Extracts were prepared from all the cells and DNA binding by STAT proteins was analysed as described in the legends to Figure 5. (b) MDA-MD-468 cells growing in full medium (FM) were treated with AG490 for 24 hours at the concentrations indicated. DNA binding by STAT proteins was analysed as described in the legend to Figure 5. STAT1 homodimers (1:1), STAT3 homodimers (3:3), heterogeneous STAT3 complexes (3H) and supershifted STAT3 complexes (3SS) are indicated.

Figure 9 shows:

Inhibition of BC cell growth. (a) Equal numbers of MDA-MB-468 and MCF-7 cells, transfected with a control vector or expression vectors for wild type of dominant-inhibitory versions of STAT3 (Y/F and E/V; see Materials and Methods) were plated in full medium and cultured for 96 hours. Cells were then harvested and counted. Data are expressed as means \pm S.D. Significant differences between groups were determined by Student's t-test. *P* values <0.05 (*) are considered significant. ** indicates *P* values <0.01 . (b) Equal numbers (1×10^5) of MDA-MB-468 and BR293 cells were plated in full medium and cultured for 60 hours. Ag490 (100 μ M) was added 48 and 24 hours prior to counting or omitted entirely. (c) As in (b) except that 1×10^6 MDA-MB-468 cells but only 5×10^5 BR293 cells were plated and the ErbB1 inhibitor PD153035 (100nM) was used. (d) As in (c) except that the irreversible ErbB1 inhibitor PD 168393 (2 μ M) was used. Inset shows ErbB1 tyrosine phosphorylation levels in MDA-MB-468 cells treated with PD 168393 (2 μ M) over 24 hours. Results are expressed as cell number after 48 hours growth, whereby error bars show the standard error from triplicate points.

Figure 10 shows: The effect of OddHL on the serum-induced accumulation of STAT1 and STAT3.

Figure 11 upper panel shows: The effect of OddHL on serum-induced phosphorylation of STAT3; middle panel the effect of N-acyl homoserine lactones (AHL) on TPA stimulation of ERKs; lower panel the effect of reactive oxygen (ROS) scavengers, and the JAK inhibitor AG490 on serum stimulation of STATs.

Figure 12 shows: Dose-response of OddHL on DNA binding by STAT1 + STAT3, inhibition of ROS-induced DNA binding by STAT1+STAT3 but not CM-induced DNA binding by STAT3.

Figure 13 upper panel shows: The effect of transcription inhibitors on serum-mediated stimulation of STAT1 + STAT3; lower panel shows that the conditioned medium from serum + OdDHL-treated cells lacks the autocrine factor required to induce STAT3 phosphorylation.

- 5 Figure 14 shows: The effect of OdDHL on breast cancer cell proliferation (right hand panels) and apoptosis (left hand panels).

Figure 15 shows:

- Serum-stimulation of STAT3 is inhibited by OdDHL. (a) BR293 cells were serum-starved (lane 1) or starved and stimulated with 10% FCS for 2 hours alone (lane 2) or in the presence of increasing concentrations of OdDHL (lanes 3-6) or 100 μ M OHHL (lane 7). Cell lysates were prepared and analysed by Western blotting for STAT3 phosphorylation with an antibody specific for STAT3 phosphorylated on Y705 (upper panel) and for STAT3 content with an antibody for STAT3 (lower panel). (b) From BR293 cells treated as in (a) nuclear extracts were prepared and analysed for STAT1 and STAT3 DNA-binding activity by EMSA with a radio-labelled probe corresponding to the M67 SIE. The inclusion of an anti-STAT3 antibody in duplicate binding reactions (even lanes) identifies STAT3-containing homo- (3:3) and heterodimers (3:1), as indicated to the right of the panel. (c) MDA-MB-468 cells were treated and analysed exactly as described for BR293 cells in (a). (d) MDA-MB-468 cells were treated and analysed exactly as described for BR293 cells in (b).

Figure 16 shows:

- OdDHL inhibits proliferation of BC cells (a) BR293 cells (1×10^6) were cultured in full medium in the presence of DMSO control (upper panel), 100 μ M OHHL (middle panel) or 100 μ M OdDHL (lower panel). After 48 hours cells were photographed. (b) Equal numbers of MCF-10F, MCF-7, MDA-MB-468 (1.5×10^6) and BR293 cells (1×10^6) were cultured in full medium alone or in the presence of various concentrations of OdDHL. After 48 hours cells were harvested and counted in a haemocytometer. Values are given as means of triplicate points, whereby error bars indicate standard errors. (c) Equal numbers of HEK293 and COS-1 cells (1×10^6) were cultured in full medium alone or in the presence of various

concentrations of OdDHL or 100 μ M OHHL. After 48 hours cells were harvested and counted in a haemocytometer. Values are given as means of triplicate points, whereby error bars indicate standard errors.

5 Figure 17 shows:

OdDHL induces apoptosis of BC cells (a) MCF-7 cells were cultured in full medium alone or in the presence of 100 μ M OdDHL or 100 μ M OHHL. After 18 hours cells were fixed, stained with DAPI and examined by confocal microscopy. Left-hand panels show DAPI, right-hand panels show the corresponding phase contrast and middle panels the image
10 overlays. (b) MDA-MB-468 cells were cultured in full medium alone or in the presence of 100 μ M OdDHL or 100 μ M OHHL. After 18 hours cells were fixed, stained and examined as in (a). (c) BR293, MCF-7 and MDA-MD-468 cells were cultured in full medium alone (lanes 1, 3, 5) or in the presence of 100 μ M OdDHL (lanes 2, 4, 6) or 400 μ M etoposide (lane 7). After 18 hours cells were lysed and analysed for PARP cleavage by Western blotting
15 with an anti-PARP antibody. Full length PARP is labelled and the lower arrow indicates the major caspase cleavage fragment.

Figure 18 shows:

OdDHL blocks the autocrine release of mitogens from BC cells (a) Equal numbers of BR293
20 cells were cultured in MEM alone, serum-free MEM conditioned by serum-stimulated BR293 cells for 2 hours, or the same supplemented with 5% FCS, as indicated. After 24 hours [3 H]-thymidine was added to the medium. After a further 18 hours, cells were harvested and analysed for incorporation of 3 H. Error bars denote SD (n = 4). (b) BR293 cells were serum-starved (lane 1) or starved and stimulated with 10% FCS for 2 hours directly (lane 2) or after
25 pretreatment with α -amanitin (100 μ g/ml) for 2.5 hours (lane 3) or actinomycin D (10 μ g/ml) for 10 min (lane 4). Nuclear extracts were prepared and analysed for STAT1 and STAT3 complexes as described in the legend to figure 1b. (c) BR293 cells were serum-starved (lane 1) or starved and stimulated with 10% FCS alone for 2 hours (lane 2), with serum-free CM alone for 15 min (lane 5) or in the presence of increasing 100 μ M AG490 (lanes 3 and 6) or
30 100 μ M OdDHL (lanes 4 and 7). Cell lysates were prepared and analysed by Western

blotting as described in the legend to figure 1a. (d) BR293 cells were stimulated with 10% FCS in the presence of 100µM OHHL (lane 1) or OdDHL (lane 2). After 2 hours the cells were washed and incubated for a further 2 hours in serum-free MEM. The conditioned medium (CM) was then used to stimulate fresh serum-starved cells for 15 min, as indicated.

5 Lysates were prepared from all the cells and analysed by Western blotting as described in the legend to figure 1a.

Figure 19 shows:

ERKs are unaffected by OdDHL (a) BR293 cells were serum-starved (lane 1) or starved and stimulated with TPA for 30 min alone (lane 2) or in the presence of increasing concentrations of OdDHL (lanes 3-6) or 100µM OHHL (lane 7). Cell lysates were prepared and analysed by Western blotting for ERK phosphorylation with an antibody specific for phospho-ERK1/2 (upper panel) and for ERK content with an antibody against ERKs (lower panel). (b) BR293 cells were serum-starved (lane 1) or starved and stimulated with Anisomycin for 30 min alone (lane 2) or in the presence of increasing concentrations of OdDHL (lanes 3-6) or 100µM OHHL (lane 7). Cell lysates were prepared and analysed by Western blotting for JNK/SAPK phosphorylation with an antibody specific for phospho-JNKs (upper panel) and for SAPK/JNK content with an antibody against JNKs (lower panel). (c) BR293 cells were treated as described in (b) and lysates were analysed by Western blotting for p38^{MAPK} phosphorylation with an antibody specific for phospho-p38 (upper panel) and for p38 content with an antibody against p38 (lower panel).

Figure 20 shows:

OdDHL potentiates STAT3 activation by EGF (a) MDA-MB-468 cells were serum-starved (lanes 1-3) or starved and stimulated with EGF for 15 min (lanes 3-6) alone or in the presence of OdDHL at 10µM (lanes 2 and 5) or 100µM (lanes 3 and 6). EGF-R was immunoprecipitated from cell lysates and analysed by Western blotting tyrosine phosphorylation with an anti phosphotyrosine antibody (upper panel, top) and EGF-R content with an anti-EGF-R antibody (lower panel, top). Cell lysates were analysed in parallel for STAT3 phosphorylation with an antibody specific for STAT3 phosphorylated on

Y705 (upper panel, bottom) and for STAT3 content with an antibody for STAT3 (lower panel, bottom). (b) Nuclear extracts were prepared from serum-starved MDA-MB-468 cells (lanes 1 and 2) or starved cells treated with EGF alone (lanes 3 and 4) or EGF and 100 μ M OHHL (lanes 5 and 6) or EGF and 100 μ M OdDHL (lanes 7 and 8), and analysed for STAT1 and STAT3 DNA-binding activity by EMSA with a radio-labelled probe corresponding to the M67 SIE. The inclusion of an anti-STAT3 antibody in duplicate binding reactions (even lanes) identifies STAT3-containing heterodimers (3:1), as indicated to the right of the panel. (c) COS-1 cells were transfected with an expression vector for STAT3, a STAT3-responsive luciferase reporter (SIE2-luc) and a control gene for β -galactosidase. After recovery and culture in starvation medium for 18 hours, cells were stimulated with EGF alone or together with AHLs or specific tyrosine kinase inhibitors, as indicated. After 6 hours cells were harvested and reporter gene expression analysed. Results are normalised against β -galactosidase values and expressed as averages \pm SD (n =3).

Figure 21 shows: In the top section: BR293, MCF-7 and MDA-MB-468 cells were serum-starved (lanes 1, 6 and 11) or starved and stimulated with 10% FCS for 2 hours, either alone (lanes 2, 7 and 12) or in the presence of 200nM Wortmannin (lanes 3, 8, 13) 100 μ M OdDHL (lanes 4, 9, 14) or OHHL (lanes 5, 10, 15). Cell lysates were then prepared and analysed by Western blotting for Akt/PKB phosphorylation with an antibody specific for Akt/PKB phosphorylated on S473 (p-Akt, upper panel) and subsequently for total Akt/PKB content with an antibody for Akt/PKB (lower panel).

In the bottom section of figure 21, BR293 cells were serum-starved (lane 1) or starved and stimulated with 10% FCS for 2 hours alone (lane 2) or in the presence of increasing concentrations of OdDHL (lanes 3-6) or 100 μ M OHHL (lane 7) or 200nM Wortmannin (lane 8). Cell lysates were prepared and analysed by Western blotting for Akt/PKB phosphorylation with an antibody specific for Akt/PKB phosphorylated on S473 (upper panel) and subsequently for total Akt/PKB content with an antibody for Akt/PKB (lower panel).

In this first section of disclosure and exemplification, the materials and methods used were as described in the following section entitled "MATERIALS AND METHODS – 1".

A number of abbreviations are used in this disclosure that are well-known and obvious to those skilled in the art. The following abbreviations are also well-known, but for clarity are defined here:

BC	Breast Carcinoma
JAK	Janus kinase
OdDHL	N-(3-oxo-dodecanoyl)-L-homoserine lactone
OHHL	N-(3-oxohexanoyl)-L-homoserine lactone
AHL	N-acyl-L-homoserine lactone
EGF	Epidermal Growth Factor

ErbB and STAT protein expression in BC cell lines

Initially, the expression levels of ErbB proteins in six BC-derived cell lines were compared by immunoblotting. As shown in Figure 2 (upper panel), ErbB1 was strongly expressed in MDA-MB-468 cells, moderately expressed in BT20 cells, weakly expressed in MDA-MB-231 cells and undetectable in the other three cell lines (MCF-7, T47D and BR293). However, MCF-7 and T47D cells have been shown previously to express low levels of surface ErbB1, indicating that the limit of detection must lie above 10,000 receptors per cell. ErbB2 was expressed at a similar level in all of the cell lines, with the exception of MDA-MB-468, in which it was undetectable. Expression of ErbB3 was also analysed and found to be moderate in MCF-7 and T47D, weak in BT20 and MDA-MB-468 and absent from MDA-MD-231 and BR293 cells. In contrast, the expression of STAT1 and STAT3 proteins in these cells showed much less variation (Figure 2, lower panel). BR293 cells alone express low levels of STAT1 proteins (lane 6). Both isoforms of STAT3 (STAT3 α and β) are expressed in all the cell lines but the β isoform is expressed at a lower level in BT20 and BR293 cells (lanes 1 and 6). Thus, these six BC-derived cell lines exhibit five different profiles of ErbB

expression, whereby only those exhibited by MCF-7 and T47D cells are similar. However, they express comparable levels of STAT1 and STAT3 proteins.

Tyrosine phosphorylation of ErbB proteins in BC cell lines

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The activity of ErbB proteins is a consequence of their tyrosine phosphorylation status. Accordingly, tyrosine phosphorylation of ErbB proteins was analysed, in those cells in which they could be detected (Figure 3), by immunoprecipitation and subsequent detection with a phosphotyrosine-specific antibody (PY20). In BT20, MDA-MB-231 and MDA-MB-468 cells, tyrosine phosphorylation of ErbB1 is weak or undetectable in normally growing cells (Figure 3, upper panel), but, as expected, it is induced (5.9, 10.8 and 8.3 fold, respectively) upon treatment of cells with EGF.

Tyrosine phosphorylation of ErbB2 is detectable in normally growing MCF-7 and T47D cells but not in the other cell lines. In MDA-MB-231 cells, EGF treatment does not elicit an increase in ErbB2 tyrosine phosphorylation, even though ErbB1 is expressed (see Figure 3) and becomes phosphorylated itself. However, in T47D and BR293 cells, which both lack ErbB1 (see Figure 3), stimulation of ErbB2 tyrosine phosphorylation by EGF is apparent (5.7 and 3.2 fold respectively).

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ErbB3 tyrosine phosphorylation is also observed under normal growth conditions in all four cell lines in which it is expressed. Moreover, in those cell lines in which ErbB1 is co-expressed, tyrosine phosphorylation of ErbB3 is induced by EGF (6.6 and 6.3 fold). In summary, although the variations in ErbB protein expression among the cell lines precludes direct quantitative comparison, those cells expressing ErbB1 display low levels of tyrosine phosphorylation on ErbB2 and ErbB3 proteins that become elevated following stimulation by EGF. Conversely, cells lines that lack ErbB1 show constitutive levels of tyrosine phosphorylation on ErbB2 and ErbB3 that remain unchanged or increase only marginally when cells are treated with EGF.

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STAT activation in BC cell lines

The phosphorylation of STAT1 and STAT3 proteins was also examined in all six cell lines with phospho-specific antibodies for each protein. As shown in Figure 4 (upper panel),
5 tyrosine phosphorylation of STAT1 was undetectable in serum-starved cells, but was stimulated in BT20 and MDA-MB-468 cells following EGF treatment (lanes 2 and 10). As already seen in Figure 2, BR293 cells express low levels of STAT1. A low level of STAT3 tyrosine phosphorylation could be seen in serum-starved BR293 cells (lower panel, lane 11), while in STAT3 immunoprecipitates probed with an anti-phosphotyrosine antibody
10 phosphorylated STAT3 was detected in all six cell lines (result not shown). Following EGF stimulation, however, tyrosine phosphorylation of STAT3 also increased in BT20 and MDA-MB-468 cells (lanes 2 and 10), mirroring the behaviour of STAT1. Because EGF-induced tyrosine phosphorylation of ErbB1 also occurs in MDA-MB-231 cells (see Figure 3), the failure to induce STAT3 tyrosine phosphorylation is likely to be a consequence of the lower
15 level of ErbB1 expression in this cell line (see Figure 2).

The function of STAT proteins depends on their DNA-binding ability, for which tyrosine phosphorylation and dimerisation are prerequisites. Initially, whole cell extracts prepared from BC cells were analysed for STAT binding activity with a cognate binding element derived
20 from the *c-fos* SIE. In extracts of serum-starved MDA-MB-468 cells, in which ErbB1 is highly expressed, a low level of heterogeneous DNA binding was detected (Figure 5, lane 1), which could be attributed, by supershift assay with anti-STAT antibodies, to STAT3 (lane 3). Control experiments confirmed that the anti-STAT3 antibody does not generate the supershifted complex (3SS), seen here and in subsequent figures, in the absence of DNA-
25 binding by STAT3 (data now shown). After stimulation of the cells with EGF, DNA-binding was much enhanced and several additional complexes were detected (lane 4) that contained STAT1 and STAT3, as evidenced by supershift assay with specific antibodies (lanes 5 and 6). In parallel experiments with BT20 cells, which also express ErbB1, EGF induced the formation of a similar set of complexes (Figure 5, lanes 10-12). However, under the same
30 experimental conditions, STAT complex formation was weaker, which may reflect the lower

ErbB1 expression in these cells (Figure 2). Furthermore, we did not detect the induction of STAT complexes by EGF in MDA-MB-231 cells, which express even less ErbB1 (result not shown). When this experiment was carried out with cells lacking ErbB1 (BR293), weak DNA binding by STAT3 was again detected in extracts of serum-starved cells, but EGF failed to stimulate the formation of additional STAT-DNA complexes (Figure 5, lanes 16-18). Thus, acute stimulation of STAT1 and STAT3 DNA-binding activity in response to EGF correlates directly with ErbB1 expression in BC cells.

Acute STAT activation requires ErbB1 and JAK kinase activity

To confirm that the acute activation of STAT DNA-binding in response to EGF was dependent upon ErbB1 kinase activity. EGF stimulation was repeated in the presence of the quinazoline inhibitor PD 153035. Pre-treatment of MDA-MB-468 cells with 100nM PD 153035 for 30 minutes inhibited tyrosine phosphorylation of ErbB1 (Figure 6a) and abrogated the induction of SIE-bound STAT complexes by EGF (Figure 6b). However, PD 153035 had no effect on the weak, heterogeneous DNA-binding by STAT3 detected by supershift assay in extracts from serum-starved cells (lane 3 and lanes 9, 12, 15, 18, 21). Thus the acute activation of STAT DNA-binding by EGF requires ErbB1 kinase activity.

The acute induction of STAT DNA-binding activity was examined in cells treated with the JAK inhibitor AG490 (34). As shown in Figure 6c, 100μM AG490 abolished STAT activation by EGF in BT20 and MDA-MB-468 cells. The weak, STAT3 DNA-binding was again unaffected (compare lane 2 with lane 6 and lane 7 with lane 9). Thus, acute stimulation of STAT DNA-binding requires both ErbB1 and JAK kinase activity, whereas the weak STAT3 DNA-binding requires the activity of neither.

A basal level of STAT3 DNA-binding similar to that in BT20 cells was also seen in serum-starved MDA-MB-231 and BR293 cells (Figure 6d) and could be detected in MCF-7 and T47D cells (not shown). In all cases, the complexes persisted in cells from which serum had been withdrawn for up to three days (Figure 6d, lane 6 and results not shown).

Serum induces elevated STAT3 activity via an autocrine signal

When serum-starved BR293 cells, which lack ErbB1, were returned to full medium, an
5 increase in STAT3 tyrosine phosphorylation over a 2 hour time course was observed (Figure
7a, upper panels). Tyrosine phosphorylation of JAK2 was also stimulated by serum over the
same period (lower panels) and both effects were blocked by AG490. As shown in Figure 7b,
STAT DNA-binding activity in nuclear (left panel) and whole cell extracts (right panel) also
10 increased, reaching a peak at 2 hours. Importantly, the STAT complexes observed in whole
cell extracts were also present in nuclear extracts, with the exception of the STAT3
complexes detected in unstimulated cells.

Compared to the rapid, acute induction by EGF, the kinetics of STAT activation in response
to serum was delayed, indicating that the upregulation of STAT DNA-binding by serum
15 involves the autocrine/paracrine mechanism now claimed.

Serum-starved BR293 cells were stimulated with 10% foetal calf serum (FCS) and, after 2
hours, half the cells were harvested while the other cells were washed thoroughly and
incubated for a further 4 hours in serum-free medium. This medium was then transferred to
20 fresh, serum-starved BR293 cells, which were incubated for a further 15 minutes. Nuclear
extracts were made from all the cells and analysed for STAT DNA-binding. As shown in
Figure 7c, STAT1 and STAT3 DNA-binding was stimulated after 2 hours by 10% FCS (lanes
3 and 4). In contrast, serum-free conditioned medium from cells incubated previously with
10% FCS for 2 hours stimulated STAT3 DNA-binding after 15 minutes (lanes 5 and 6).
25 Similarly, conditioned medium from MDA-MD-468 cells cultured for 2 hours with 10% FCS
was able to stimulate STAT3 DNA-binding in BR293 cells within 15 minutes (result not
shown). Treatment of BR293 cells with conditioned medium also induced tyrosine
phosphorylation of STAT3 within 15 minutes, whereas EGF treatment did not (Figure 7d).
Demonstrating that BC cells cultured in 10% FCS release factors that stimulate tyrosine
30 phosphorylation of STAT3 and its consequent DNA-binding activity.

As BR293 cells do not express ErbB1, the involvement of ErbB1 in the serum-dependent activation of STAT3 is unlikely. Consistent with this inference, when FCS was applied to serum-starved MDA-MB-468 cells pre-treated with PD 153035, the delayed serum stimulation of STAT3 DNA-binding was not affected (result not shown). The role of JAKs in the serum-dependent activation of STAT3 was further assessed by pre-treating BR293 cells with 100 μ M AG490 for 30 minutes which completely blocked STAT3 activation (Figure 8a, lanes 5 and 6). However, when conditioned medium from BR293 cells was applied to serum-starved cells treated with AG490, no inhibition was observed (lanes 17 and 18). Showing that primary signal mediating STAT3 activation by serum requires JAK activity, whereas the secondary autocrine signal acts independently of JAKs. MDA-MB-468 cells growing in full medium were treated with AG490 for 24 hours, as shown in Figure 8b, this reduced the elevated, serum-dependent level of STAT3 activity to a constitutive basal level.

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Taken together, the preceding results distinguish three distinct levels of STAT activity in BC cells, as manifested by DNA-binding. Firstly, in cells expressing ErbB1, several STAT-DNA complexes can be induced acutely by EGF, which is dependent upon the kinase activity of both ErbB1 and JAKs. Secondly, an intermediate level of DNA-binding is induced by serum via an autocrine mechanism involving JAKs but not ErbB1. Thirdly, a weak, constitutive level of DNA-binding by STAT3, which is independent of ErbB1 and JAKs, is detected in whole cell extracts from serum-starved cells and persists for up to 3 days.

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Inhibition of BC cell growth

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MDA-MB-468 and MCF-7 cells were transfected with expression vectors for STAT3 and two *trans*-dominant negative mutants thereof. In each instance, the dominant negative mutants caused a 25-30% decrease in the growth of transfected cells over 4 days (Figure 9, top left panel). As only a proportion of the cells was transfected, this result probably underestimates the effect of dominant-inhibitory STAT3 mutants on BC cell growth. STAT3 is

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therefore crucial in the proliferation of these BC cell lines. The effects of JAK inhibition on BC cell growth was measured. As shown in Figure 9 (top right panel), treatment of MDA-MB-468 cells and BR293 cells with AG490 for 24 or 48 hours had a dramatic effect on cell growth, reducing cell proliferation by >75% over 48 hours. Thus, JAK function is important for BC cell proliferation. As JAKs are involved in both the acute and intermediate levels of STAT3 activity, the effect on cell growth of two specific ErbB1 inhibitors, PD 153035 and PD 168393 was also measured. Treatment of MDA-MB-468 and BR293 cells with either reagent had no effect on their proliferation over 48 hours (lower panels). To demonstrate that ErbB1 was indeed inhibited, MDA-MB-468 cells cultured and treated with PD 168393 in parallel were stimulated at different time points with EGF for 15 minutes and tyrosine phosphorylation of ErbB1 was measured. PD 168393 completely inhibited ErbB1 tyrosine kinase activity over 24 hours (Figure 9, inset). This data shows that BC cell proliferation correlates with STAT3 activity which is maintained by the serum-dependent autocrine/paracrine pathway now claimed.

OdDHL blocks serum stimulation of STAT1 and 3.

Serum-starved BR293 cells were pre-treated with 200 μ M OdDHL (active) or OHHL (inactive) for 30 min. and then stimulated with serum for 2h. Cells were lysed and the DNA-binding activity of STAT1 and 3 was examined by EMSA. As shown in figure 10 (left panel), pre-incubation of the cells with OdDHL, but not OHHL, prevented serum induction of STAT1 and 3 complexes. Treatment of serum-starved BR293 cells with either AHL alone for up to 2h did not induce STAT complex formation (right hand panels). Similarly, as shown in figure 11 (upper panel), pre-incubation of BR293 cells with OdDHL, but not OHHL or an unrelated signal molecule (PQS), blocked the serum-induced phosphorylation of STAT3. (This panel is the first part of an experiment also shown in figure 13.)

None of the three compounds had any noticeable effect on the activation of the ERK Mitogen-Activated Protein Kinase (MAPK) cascade by TPA, as indicated by their failure to prevent phosphorylation of ERK1/2 (middle panel). This indicates that the action of OdDHL

on STAT activation is not part of an unspecific, pleiotropic effect. Other small molecules also inhibited serum induction of STAT1 and STAT3 DNA-binding activity, for example scavengers of reactive oxygen species (ROS) such as N-acetyl cysteine (NAC) and diphenylene iodonium chloride (DPI) and the JAK inhibitor AG490 (see Figure11, lower panel).

OdDHL blocks autocrine factor release from BC cells

A titration of OdDHL revealed that its IC50 for the inhibition of serum-induced STAT1 and STAT3 DNA-binding activity lies between 50-100µM (Figure 12, left hand panel). OdDHL, but not OHHL, also blocks the activation of STAT1 and STAT3 DNA binding by H₂O₂, which increases intracellular ROS, but not the activation of STAT 3 DNA binding by conditioned medium (CM) from BR293 cells (right hand panels). This indicates that OdDHL inhibits the STAT3 activation pathway upstream of autocrine factor release (see Figure 1).

Autocrine factor release is frequently associated with *de novo* gene expression and protein synthesis. As shown in figure 13 (upper panel), both α -amanitin and Actinomycin D (inhibitors of transcription) reduce the levels of STAT1 and STAT3 activation by serum whereas methanol (the vehicle) has no inhibitory effect, indicating that this pathway involves gene expression.

If OdDHL can block autocrine factor release, then conditioned medium from treated cells should not elicit STAT3 activation. Indeed, CM from serum-stimulated BR293 cells pre-treated with OdDHL was unable to stimulate STAT3 phosphorylation, whereas CM from serum-stimulated cells pre-treated with OHHL or PQS could do so (Figure 13, lower panels).

OdDHL blocks BC cell oroliferation and induces apoptosis. BR293 cells were grown in full medium (10% FCS) in the absence or presence of OdDHL or the control OHHL. After 24h, cell were harvested, stained with DAPI and examined by fluorescence microscopy. OdDHL induced DNA condensation and nuclear fragmentation whereas untreated or OHHL-treated cells remained viable (figure 14, left hand panels). The effect of OdDHL on proliferation

was also apparent: cells cultured in the presence of OdDHL for 48h grew poorly or not at all, but control cells plated at the same density reached confluence (right hand panels).

Taken together, these results show that serum-dependent STAT3 activity is required for BC
5 cell proliferation and that biologically active AHLs such as OdDHL block activation of
STAT1 and 3, thereby inhibiting BC cell proliferation.

In the following section of description and exemplification, the materials and methods
referred to are described fully in the section entitled "Materials and Methods - 2" that
10 follows.

The human pathogen *Pseudomonas aeruginosa* uses quorum-sensing signal molecules
(QSSMs) to regulate virulence gene expression. It has been shown that such molecules are
also able to suppress host immune responses of the type commonly associated with auto-
15 immune disease, although the mechanism of action is obscure. However, regulation of
immune function is known to involve STAT proteins.

Here we have explored the possibility that QSSMs of *P. aeruginosa* are able to modulate
STAT activity in the context of BC cell growth. We show that constitutive STAT3 activity in
proliferating human BC cells is down-regulated by the QSSM *N*-(3-oxododecanoyl)-L-
20 homoserine lactone OdDHL, resulting in apoptotic cell death. These results support the
notion of OdDHL as a bioactive molecule in eukaryotic systems and as a paradigm for a
novel class of antiproliferative molecules.

T cell responses to immune challenge are orchestrated by a complex array of cytokines with
25 diverse and often selective effects on their target cells, controlling, among other things, cell
survival and proliferation. Inextricably linked to cytokine action are intracellular signalling
pathways that involve Signal Transducer and Activator of Transcription (STAT) proteins and
a family of protein tyrosine kinases referred to as Janus Kinases (JAKs) (20). Activated
cytokine receptors provide scaffolds upon which STATs are phosphorylated by JAKs,

whereupon STATs translocate to the nucleus and up-regulate the expression of target genes, which include genes for numerous cytokines (10, 17).

The intimate associations between pathogen and host make it highly likely that the former have evolved subtle and selective strategies to subvert the immune systems of the latter.

- 5 Notably, the widely used immune-suppressant drugs Cyclosporin A and Rapamycin are both naturally occurring compounds. In this context it has been shown recently that quorum-sensing signal molecules (QSSMs) from *Pseudomonas aeruginosa* are able to suppress immune responses of the type commonly associated with auto-immune disease (25). In contrast, more recent data suggests that these molecules may have pro-inflammatory activity
- 10 (23). However, the mechanism underlying these responses is obscure and the molecular target for OdDHL remains to be identified. Given their involvement in cytokine-mediated events, elements of the JAK/STAT pathway would appear to offer prime targets for pathogens aiming to evade or inactivate the immune systems of their hosts.

- STAT proteins are implicated in cellular processes distinct from those regulating the immune
- 15 system. For example, STAT3 plays a role in driving cell proliferation and counteracting differentiation signals (2), while a STAT3 mutant that dimerises in the absence of tyrosine phosphorylation is constitutively active and functions as an oncogene (4). Moreover, the proliferation of a range of tumour-derived cells, notably of Breast Carcinoma (BC) origin, has been shown by several groups to depend on the constitutive activity of STAT3 (5, 7, 11).

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- Given the importance of the JAK/STAT pathway for immune function and of STAT3 for cell proliferation, we decided to explore the possibility that AHLs might modulate STAT signalling in the context of cell proliferation. Here we show that STAT1 and STAT3 activities in proliferating human BC cells are down-regulated by *N*-(3-oxododecanoyl)-L-homoserine lactone (OdDHL), the major QSSM of *P. aeruginosa* (27) (16), resulting in
- 25 apoptotic cell death. However, in cells stimulated by EGF the acute activation of STAT3 is augmented by OdDHL. Our findings indicate that OdDHL is a bioactive molecule in eukaryotic systems and a paradigm for a novel class of antiproliferative molecules. They also

raise the possibility that in order to serve disparate roles STAT3 may be partitioned into two functional populations, whereby disruption of one automatically augments the other.

Inhibition of STAT activity by bacterial QSSMs

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As STAT3 activity appears to be important for BC cell proliferation, it was of interest to identify inhibitors of STAT function and test their consequent effects on BC cell growth. Among the potential inhibitors chosen for analysis were bacterial QSSMs, some of which have recently been shown to influence aspects of host immune function, in which the JAK/STAT signalling pathway is notionally involved (25). Pretreatment of BR293 cells for 15 minutes with increasing concentrations of OdDHL inhibited serum-induced STAT3 tyrosine phosphorylation at 100 μ M (Figure 15a, lane 6). In contrast, the short chain analogue *N*-butanoyl homoserine lactone (OHHL), which lacks immune-modulatory activity, had only a slight effect at the same concentration (lane 7). Similar results were obtained with MDA-MB-468 cells (figure 15a, lower panel).

The effects of AHLs on STAT3 DNA binding were also monitored in parallel. Nuclear extracts were prepared and assayed with a high affinity STAT1/STAT3 binding site (M67 SIE 9 (29)) and STAT3-containing complexes were identified by including a STAT3-specific antibody in duplicate binding reactions. In this assay OdDHL was seen to inhibit DNA binding by STAT3 at 50 μ M (Figure 16b, lanes 9 and 10), while again OHHL had only a marginal effect at 100 μ M (lanes 13 and 14). The lower IC₅₀ for DNA-binding suggests that the negative impact of OdDHL on STAT phosphorylation may be an indirect consequence of OdDHL acting primarily on a subsequent event in the STAT activation mechanism. Again, similar results were obtained with MDA-MB-468 cells, except that only the STAT3 homodimer was detected (figure 15b, lower panel).

It is noteworthy that the small increase in STAT3 phosphorylation at 50 μ M OdDHL is observed consistently in BR293 cells, as is the increase in DNA binding by STAT1 and STAT3 at 20 μ M. In MDA-MB-468 cells, an increase is also observed and is apparent at

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10µM OdDHL. This may be explained by OdDHL having opposite effects on two signal pathways converging on STATs (see also below). In summary, OdDHL, but not OHHL, severely impairs STAT3 activation in serum-stimulated cells.

5 OdDHL inhibits BC cell proliferation

Treatment of BC cells with OdDHL had a marked effect on their proliferation. As shown in figure 16a, BR293 cells treated with vehicle (DMSO) or OHHL reached confluence in 48 hours, whereas those exposed to OdDHL grew poorly or not at all. Indeed, OdDHL markedly
10 inhibited the proliferation of three tumorigenic BC cell lines (MCF-7, BR293 and MDA-MB-468) by 70%-80% over a 48-hour period. However, its effect on the proliferation of non-tumorigenic breast epithelial cells (MCF-10F), which were previously shown to be insensitive to the JAK inhibitor AG490 (5, 11), was slight (Figure 16b). We also examined the effects of OdDHL and OHHL on transformed human and simian cell lines. Growth of
15 HEK293 cells, which have constitutive STAT3 activity, was sensitive to treatment with OdDHL, whereas proliferation of COS1 cells, which are known to express low levels of STAT proteins (22), was unaffected (Figure 16c). These findings again link constitutive STAT3 activity to cell proliferation.

20 OdDHL induces apoptosis in proliferating BC cells

The low numbers of BC cells surviving for 48 hours in the presence of OdDHL suggested that they might be undergoing apoptotic cell death. To assess this possibility, cell integrity was analysed. In growing MCF-7 and MDA-MB-468 cells treated with OdDHL for 24 hours, staining with 4',6'-diamidino-2-phenylindole (DAPI) revealed evidence of nuclear
25 disruption, possibly indicative of apoptosis, whereas control cells or cells treated similarly with OHHL were unaffected (Figure 17a). Therefore the integrity of poly ADP-ribose polymerase (PARP), a well-characterised target for caspases, was also examined. Although BR293 cells express low levels of PARP, the characteristic cleavage fragment was detected in cells treated with OdDHL (Figure 17b, lane 2). OdDHL also induced PARP cleavage in
30 MCF-7 cells (lane 4) and in MDA-MB-468 cells to the same extent as etoposide (compare

lanes 6 and 7). Taken together, these results indicate that STAT3 inhibition by OdDHL causes apoptosis in proliferating BC cells.

OdDHL downregulates autocrine release from BC cells

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Serum has been shown to stimulate the release of an autocrine factor(s) that contributes to STAT3 activation in BC cells (11). Autocrine secretion of Prolactin has previously been reported to activate JAK2 and hence ErbB2 in BC cells (28). In addition, angiotensin II was found to stimulate autocrine release of IL-6 from rat cardiomyocytes, resulting in elevated STAT activity (19). However, as discussed previously (11), several criteria appear to distinguish these mechanisms from the autocrine-mediated STAT3 activation pathway in BC cells.

As shown in figure 18a, BR293 cells cultured in low serum undergo DNA synthesis, as measured by ³H-thymidine incorporation. However, when CM was supplemented with 5% serum, the level of DNA synthesis doubled, indicating that CM contains one or more mitogens released by BR293 cells.

Given the time delay between serum stimulation and the resultant activation of STAT3 (11), it was conceivable that the autocrine process involved *de novo* gene expression. Pretreatment of cells with α -amanitin for 2.5 hours or with Actinomycin D for 10 minutes blocked the stimulation of DNA binding by STAT1 and STAT3 (Figure 18b, lanes 3 and 4), which is consistent with this notion.

The JAK inhibitor AG490 inhibits BC cell proliferation (7) and interferes with STAT3 activation by serum but not CM (11). Similarly, pretreatment of cells with OdDHL prevented the delayed STAT3 phosphorylation in response to serum (Figures 18c, lane 4; see also Figure 15a) but not the rapid response to CM (Figure 18c, lane 7). However, as shown in Figure 18d, CM from serum-stimulated cells that had been pretreated with OdDHL failed to induce STAT3 phosphorylation (lane 4) while CM from cells pretreated with OHHL was active (lane 3), indicating that OdDHL-treated cells fail to release the active autocrine factor.

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Influence of OdDHL on MAPK cascades

The effects of OdDHL on STAT3 activation by serum prompted us to monitor its effects on other signalling pathways. One downstream consequence of serum stimulation of cells is the activation of MAPK cascades, reflected by the phosphorylation of Extracellular signal-regulated Kinases (ERKs) and, in some cases, cJun N-terminal Kinases/Stress-Activated Protein Kinases (JNK/SAPK) and p38-family MAPKs (9, 18). We therefore measured the effects of OdDHL on MAPK cascades activated by pathway-specific stimuli. Neither OdDHL nor OHHL had any effect on the activation of ERKs in response to TPA (Figure 19a) or serum (result not shown). In contrast, OdDHL, but not OHHL, caused a modest but reproducible inhibition of p46/p54 JNK/SAPK (Figure 19b) and p38^{MAPK} phosphorylation (Figure 19c) in cells treated with anisomycin, a well-established stress agonist. However, it is unlikely that this inhibition of stress-activated MAPKs contributes to the anti-proliferative effects of OdDHL

OdDHL enhances STAT activation in response to EGF

OdDHL blocks STAT3 activity in proliferating BC cells and precipitates cell death by apoptosis, providing further evidence for a link between STAT3 and cell proliferation. However, it remained to be seen if OdDHL also blocks STAT3 activation in response to acute stimulation. Although BR293 cells lack the EGF receptor (EGF-R) and do not respond to EGF, MDA-MB-468 cells, which are equally susceptible to OdDHL (Figure 15), express high levels of the receptor (11). We therefore tested the influence of OdDHL on the activation of STAT3 by EGF in MDA-MB-468 cells, which express the EGF-R. As shown in Figure 20, the outcome was markedly different. OdDHL alone at 100µM caused an increase in detectable tyrosine phosphorylation of the EGF-R in unstimulated cells (Figure 20a, lane 3) and potentiated receptor tyrosine phosphorylation in response to EGF stimulation several fold, even at lower concentrations (10µM) (compare lanes 5 and 6 with lane 4). Similarly, STAT3 phosphorylation in response to EGF was augmented by co-treatment of cells with 100µM OdDHL (Figure 20a, lower panels), while in DNA-binding assays

activation of STAT3 was seen to be enhanced by OdDHL but unaffected by OHHL (Figure 20b, compare lanes 7 and 8 with lanes 3-6). Moreover, in COS-1 cells transfected with an expression vector for murine STAT3, the 5-6 fold stimulation of STAT3-dependent reporter gene expression by EGF was augmented by co-treatment of cells with OdDHL (Figure 20c).

- 5 In contrast, OHHL had no effect (not shown). The involvement of the EGF-R and JAKs in these events was confirmed with the inhibitors PD153035 and AG490 respectively. Thus, in stark contrast to its effects on serum-dependent STAT3 activity, OdDHL has a positive effect on STAT3 activation in response to EGF.

10 DISCUSSION

- STAT proteins are fundamentally involved in implementing the changes in gene expression that coordinate numerous biological programmes, such as haematopoiesis, embryogenesis and immune responses. STAT3 in particular has also been linked to cell proliferation and survival and shown to possess oncogenic potential (2, 3). Based on the
- 15 premise that pathogens gain advantage by subverting host immune systems, in which STATs play pivotal roles, we explored the possibility that QSSMs of *P. aeruginosa* can modulate STAT activity. We found that OdDHL, but not OHHL, is able to potentiate acute stimulation of STAT3 by EGF, but to down-regulate STAT3 activity and induce apoptosis in the context of proliferating BC cells.

20

Role of STAT3 in BC cell proliferation

- A large body of evidence implicating STAT3 as a positive regulator of proliferation in a range of tumour tissues has accumulated. Evidence for the role of STAT3 included the effects
- 25 of dominant-inhibitory STAT3 mutants, which were found to reduce proliferation of several tumour cell types, counteract transformation by several oncogenes and exert a negative-selection on the establishment of stable cell lines in which they were expressed (reviews in (26).

- STAT3 is involved in the expression of several proteins that participate in cell cycle control. It appears to mediate the induction of *c-myc* in response to growth factors including IL-6 and various oncogenes including *v-src* and *v-abl* (reviewed in (8), which would contribute to G₁-S progression. In addition, *Pim1* and *Pim2* have been identified as STAT3-responsive genes.
- 5 *Pim1* encodes a serine/threonine kinase that phosphorylates and activates Cdc25A, a major regulator of cyclin-dependent kinases. In the absence of STAT3 activity constitutive expression of both *Pim1* and *c-Myc* was shown to be required for cell cycle progression (14, 21).
- 10 STAT3 can also influence the balance between survival and apoptotic signals. Several lines of evidence indicate that pro-survival Bcl-family members are upregulated by STAT3. For example the high levels of Bcl-x_L expressed in the Fas-resistant myeloma cell line U266 are reduced upon STAT3 inactivation, whereupon the cells undergo apoptosis (6). These findings highlight the role of STAT3 as a positive regulator of the cell cycle and anti-
- 15 apoptotic signalling in at least a subset of human cell types.

Distinct pathways for STAT3 activation

- The most intriguing of our findings relates to the converse effects of OdDHL on two distinct
- 20 modes of STAT3 activation. Whereas the serum-dependent STAT3 activity was abrogated by OdDHL, acute activation of STAT3 in response to EGF was enhanced, an effect apparent at the level of EGF-R phosphorylation. STAT3 activation in response to EGF is known to require the kinase activities of its receptor and JAK2, whereas serum stimulation involves only the latter (11). OdDHL is therefore unlikely to affect JAK2 directly and indeed,
- 25 preliminary data suggest that OdDHL does not interfere with JAK2 phosphorylation in serum-stimulated cells (LL and PES, unpublished). JAK signalling appears to extend beyond STAT activation as tyrosine residues phosphorylated by JAK2 on either ErbB2 or gp130 have been shown to serve as docking sites for Grb2 and SHP2 respectively. Both proteins, either directly or through Gab1, can transmit signals to the ERK cascade (8) and references
- 30 within (28). Activation of this hypothetical pathway would be consistent with immediate

early gene expression and resultant autocrine secretion, which are implicated as downstream events in the pathway blocked by OdDHL.

One possible explanation for the reciprocal effects of OdDHL on STAT activation by EGF and serum is that augmentation of the one occurs to the detriment of the other. Several models for the activation of STATs have been proposed, a common aspect of which is the recruitment of STATs to phosphorylated receptor chains from a pool of latent cytoplasmic monomers. However, recent reports indicate that a fraction of cytoplasmic STATs is not present as monomers but rather in multi-protein complexes (15). Conceivably, distinct STAT fractions are targeted by different activation mechanisms. Following growth factor stimulation, nuclear translocation of STAT3 has been shown to require receptor-mediated endocytosis, but the existence of alternative pathways has not been ruled out (1). Notably, separate nuclear import pathways have been identified for monomers and dimers of STAT1 that are predicted to be conserved among STAT proteins (13). Thus there is scope for OdDHL to act selectively during STAT activation.

An alternative possibility is that OdDHL has two molecular targets, one that is activated, possibly at low OdDHL concentrations, and another that is inactivated at higher concentrations. This model would be consistent with the observed increase in STAT3 phosphorylation in response to serum at lower OdDHL concentrations and the sharp threshold for inactivation above 50 μ M seen in figure 15. The nature of serum as a mixed agonist means that multiple signals are likely to contribute to the net stimulation of STAT proteins. We are currently using a series of chemical analogues to test this possibility. Ultimately the identification of the molecular target(s) for OdDHL in eukaryotic cells will help to resolve these questions.

We analysed the effect of both OdDHL and OHHL on the Phosphatidylinositol-3OH Kinase (PI3K) survival pathway. As shown in figure 21, Akt phosphorylation was induced in serum-stimulated cells and blocked, as expected, by Wortmannin, an established inhibitor of PI3K. OdDHL blocked phosphorylation of Akt/PKB to a similar degree as Wortmannin in BR293,

MCF-7 and MDA-MB-468 cells. A titration showed it to be effective only at concentrations above 50 μ M. OHHL had no effect on PI3K signalling.

MATERIALS AND METHODS - 1

5

Cell culture and extract preparation

Breast cancer cell lines (BR293, BT20, MCF-7, MDA-MB-231, MDA-MB-468, T47D) were maintained in Minimum Essential Medium Eagle (MEM, Sigma) supplemented with 10% foetal bovine serum (FCS), 1% MEM non-essential amino acids, 1% glutamine and 1% penicillin-streptomycin at 37°C under 5% CO₂. These cell lines are well-known, and widely available to persons in the field. In particular, three of the lines are deposited with the American Type Culture Collection (ATTC) with the following identifying codes:

Cell line	ATCC Number
MDA-MB-468	HTB-132
MCF7	HTB-22
MDA-MB-231	HTB-26

15

For preparation of extracts for electrophoretic mobility shift assays (EMSAs), cells were seeded in 6-well plates (Costar) and cultured until confluent. Thereafter the cells were maintained in serum-free medium overnight before application of appropriate stimuli. For whole cell extracts, cells were lysed in TSET buffer (10mM Tris-HCl, pH7.0, 50mM NaCl, 1mM EDTA, 1% Triton X-100) supplemented with 2mM Na₃ VO₄, 5mM Na₄P₂O₇, 5mM NaF, 5mM EDTA, 2mM Benzamidine, 0.2mM PMSF, 1mM DTT and 1 μ g/ml of each leupeptin, aprotinin, pepstatin.

25 Extractions were cleared by centrifugation at 16,000 x g for 10 minutes, snap-frozen in liquid N₂ and stored at -80°C. Nuclear extracts were prepared in high salt hypertonic buffer (20mM

HEPES pH7.9, 420mM NaCl, 20% glycerol, 1mM EDTA, 1mM EGTA, 0.2% NP-40, 20mM NaF, 1mM Na₃ VO₄, 1mM Na₄P₂O₇, 2mM Benzamidine, 0.5mM PMSF, 1mM DTT and 1µg/ml each of leupeptin, aprotinin and pepstatin.

- 5 For immunoprecipitation and immunoblotting experiments, cells were grown to confluence in 10cm dishes and maintained in full medium or starved in serum-free medium overnight before the application of appropriate stimuli. Lysates were prepared in TBSN buffer (50mM Tris-HCl pH7.5, 150mM NaCl, 1mM EDTA, 1% NP-40) supplemented with protease inhibitors (1mM Na₃ VO₄, 10mM Na₄P₂O₇, 10mM NaF, 5mM EGTA, 10mM Benzamidine, 10 0.2mM PMSF and 1µg/ml each of leupeptin, aprotinin, pepstatin). Lysates were cleared by centrifugation at 16,000 x g for 10 minutes and used directly for immunoprecipitations or stored at -20°C for further use. AG490 was purchased from Sigma: PD 153035 was provided by Glaxo Wellcome and PD 168393 was purchased from Calbiochem.

15 Antibodies

- The αSTAT1, αSTAT3, αphosphotyrosine (PY20) and αErbB1 monoclonal antibodies were purchased from Transduction Laboratories; the αErbB2, αJAK2 antisera, the αErbB3 monoclonal antibody, the αphospho-STAT1 (polyclonal) and the αphospho-STAT3 20 (monoclonal) antibodies were purchased from Upstate Biotechnology; the rabbit polyclonal αSTAT1 and αSTAT3 antisera were made in our laboratory. The αErbB1 monoclonal antibody used for immunoprecipitations was kindly provided by Dr Lindy Durrant (University of Nottingham, UK).

25 Immunoprecipitation and immunoblotting

Equal amounts of lysates were incubated with the appropriate antibody for 2 hours at 4°C. Immune complexes were then allowed to bind to protein A-Sepharose beads for 1 hour at 4°C and collected by centrifugation. Immunoprecipitates were washed three times in 100mM Tris-

HCl pH 7.5, 100mM NaCl, 1mM EDTA, 0.1mM PMSF, 0.5% NP-40. Thereafter, samples were taken up in sodium dodecyl sulphate (SDS) loading buffer and boiled for 5 min.

5 Samples were separated by electrophoresis through 6% polyacrylamide-SDS gels. Proteins were then transferred to polyvinylidene difluoride (PVDF) membranes with a semi-dry electroblotting apparatus. The membranes were incubated with appropriate primary antibodies at room temperature for 1 hour or 4°C overnight according to suppliers' instructions, washed and stained with horseradish peroxidase-coupled secondary antibodies. The membranes were developed with an enhanced chemiluminescence kit (Amersham).

10

Electrophoretic Mobility Shift Assays (EMSAs)

15 DNA binding assays were carried out as previously described. Briefly, DNA binding by STAT proteins was analysed with a ³²P-labelled oligonucleotide duplex (M67SIE). Extracts were incubated with the DNA probe and protein-DNA complexes were separated by electrophoresis on 5% polyacrylamide gels containing 2.5% glycerol in 0.5 x Tris-Borate-EDTA (TBE) buffer. After separation, the gels were fixed, dried and analysed with a phosphorimager (Fuji). For supershift analyses of STAT-DNA complexes, extracts were pre-
20 incubated with αSTAT1 or αSTAT3 antisera at room temperature for 1 hour. The oligonucleotide probe was then added and the EMSA was performed as described above.

Plasmids and Oligonucleotides

25 The expression vectors for wild type and dominant-negative STAT3 proteins (STAT3-E/V) and STAT3-Y705F) were generous gifts of Drs Curt Horvath (Mount Sinai, USA) and James E Darnell Jr (Rockefeller, USA) and have been characterised previously.

The sequences of the oligonucleotides used to generate the M67 EMSA probe, which was
30 derived from the vSis-inducible element (SIE) of the human *c-fos* promoter are:

Upper strand: 5'-CTAGCATTTCCTGAAAT

Lower strand: 5'-CTAGATTACGGGAAATG

5 Cell proliferation assays

Equal numbers of BR293 and MDA-MB-468 cells were seeded in MEM containing 10% FCS into 10cm dishes. Cells were allowed to grow in the presence of the JAK inhibitor AG490 (100 μ M) or the ErbB1 inhibitors PD 153035 (100nM) and PD 168393 (2 μ M) for 24 or 48 hours. For the 48 time points, fresh medium containing the appropriate inhibitor was applied to the cells after 24 hours. Controls were allowed to grow for 48 hours in the absence of inhibitor. Thereafter, all the cells were washed twice with ice cold PBS, harvested and counted under a phase-contrast microscope. Values are expressed as averages \pm S.D. (n=3).

For the proliferation assays with wild type and dominant negative STAT3 mutants, equal numbers of MDA-MD-468 or MCF-7 cells, maintained in MEM supplemented with 10% FCS, were transfected by DNA-calcium co-precipitation with 4 μ g of the corresponding expression vector or the control vector(pRc/CMV). After 96 hours, cells were harvested and processed as described above.

MATERIALS AND METHODS – 2

Cell culture and extract preparation

Breast cancer cell lines (BR293, MCF-7 and MDA-MB-468) were maintained in Minimum Essential Medium Eagle (MEM, Sigma) supplemented with 10% foetal calf serum (FCS), 1% MEM non-essential amino acids, 1% glutamine and 1% penicillin-streptomycin at 37°C under 5% CO₂. MCF-10F cells, one of a series of non-tumorigenic lines derived from benign breast epithelial tissue, were grown as adherent cells in a 2:1 mixture of Minimum Essential Medium Eagle and Ham's F12 medium (Sigma) supplemented with 5% horse serum, 2 mM

glutamine, 10 $\mu\text{g ml}^{-1}$ insulin, 20ng ml^{-1} EGF, 100ng ml^{-1} cholera toxin, 0.5 $\mu\text{g ml}^{-1}$ hydrocortisone and 1% gentamycin. These cell lines are well-known, and widely available to persons in the field. In particular, the following lines are deposited with the American Type Culture Collection (ATCC) with the following identifying codes:

5

Cell line	ATCC Number
MDA-MB-468	HTB-132
MCF7	HTB-22
MDA-MB-231	HTB-26

For the preparation of nuclear extracts for electrophoretic mobility shift assays (EMSAs), cells were seeded in 10cm dishes and cultured until confluent. Thereafter the cells were maintained in serum-free medium overnight before application of appropriate stimuli.

- 10 Nuclear extracts were prepared as described previously (11) in high salt hypertonic buffer (20mM HEPES pH7.9, 420mM NaCl, 20% glycerol, 1mM EDTA, 1mM EGTA, 20mM NaF, 1mM Na_3VO_4 , 1mM $\text{Na}_4\text{P}_2\text{O}_7$, 2mM Benzamidine, 0.5mM PMSF, 1mM DTT and 1 $\mu\text{g/ml}$ each of leupeptin, aprotinin and pepstatin.

- 15 For immunoprecipitation and immunoblotting experiments, cells were grown to confluence in 10cm dishes and maintained in full medium or starved in serum-free medium overnight before the application of appropriate stimuli. Lysates were prepared in TBSN buffer (50mM Tris-HCl pH7.5, 150mM NaCl, 1mM EDTA, 1% NP-40) supplemented with protease inhibitors (1mM Na_3VO_4 , 10mM $\text{Na}_4\text{P}_2\text{O}_7$, 10mM NaF, 5mM EGTA, 10mM Benzamidine, 0.2mM PMSF and 1 $\mu\text{g/ml}$ each of leupeptin, aprotinin and pepstatin). Lysates were cleared by centrifugation at 16,000 x g for 10 minutes and used directly for immunoprecipitations or stored at -20°C for further use. AG490 was purchased from Sigma; PD153035 was provided by Glaxo-Smith-Kline and PD168393 was purchased from Calbiochem.

- 25 Plasmids and Oligonucleotides

The luciferase reporter construct pSIE2-luc contains 2 copies of the M67 site inserted upstream of the adenovirus 2 E4 basal promoter.

The sequences of the oligonucleotides used to generate the M67 EMSA probe, which was derived from the vSis-inducible element (SIE) of the human *c-fos* promoter, are:

Upper strand: 5'-CTAGCATTTCCTCGTAAAT

Lower strand: 5'-CTAGATTACGGGAAATG

10 Antibodies

The anti-STAT3, anti-phosphotyrosine (PY20) and anti-ErbB1 monoclonal antibodies were purchased from Transduction Laboratories; the anti-phospho-STAT3 (monoclonal) antibody was purchased from Upstate Biotechnology; the rabbit polyclonal anti-STAT3 antisera was made in our laboratory (11); the anti-ErbB1 monoclonal antibody used for immunoprecipitations was kindly provided by Dr. Lindy Durrant (University of Nottingham, UK); the anti-PARP antibody was purchased from New England Biolabs.

Immunoprecipitation and immunoblotting

20 Equal amounts of lysates were incubated with the appropriate antibody for 2 hours at 4°C. Immune complexes were then allowed to bind to protein A-Sepharose beads for 1 hour at 4°C and collected by centrifugation. Immunoprecipitates were washed three times in 10mM Tris-HCl pH 7.5, 100mM NaCl, 1mM EDTA, 0.1mM PMSF, 0.5% NP-40. Thereafter, samples were taken up in sodium dodecyl sulphate (SDS) loading buffer and boiled for 5 min.

25

Samples were separated by electrophoresis through 6% polyacrylamide-SDS gels. Proteins were then transferred to polyvinylidene difluoride (PVDF) membranes with a semi-dry electroblotting apparatus. The membranes were incubated with appropriate primary antibodies at room temperature for 1 hour or 4°C overnight according to suppliers'

instructions, washed and stained with horseradish peroxidase-coupled secondary antibodies. The membranes were developed with an enhanced chemiluminescence kit (Amersham).

Electrophoretic Mobility Shift Assays (EMSAs)

5

DNA binding assays were carried out as previously described (11). Briefly, DNA binding by STAT proteins was analysed with a ^{32}P -labelled oligonucleotide duplex (M67SIE). Extracts were incubated with the DNA probe and protein-DNA complexes were separated by electrophoresis on 5% polyacrylamide gels containing 2.5% glycerol in 0.5 x Tris-Borate-EDTA (TBE) buffer. After separation, the gels were fixed, dried and analysed with a phosphorimager (Fuji). For supershift analyses of STAT-DNA complexes, extracts were pre-incubated with anti-STAT3 antiserum at room temperature for 1 hour. The oligonucleotide probe was then added and the EMSA was performed as described above.

Cell proliferation assays

Equal numbers of BR293, MCF-10F, MCF-7, MDA-MB-468, HKEK293 and COS1 cells were seeded in the appropriate growth medium into 10cm dishes. Cells were allowed to grow in the presence of various concentrations of OdDHL or OHHL for 24 or 48 hours. For the 48-hour time points, fresh medium containing the appropriate inhibitor was applied to the cells after 24 hours. Controls were allowed to grow for 48 hours in the absence of inhibitor. Thereafter, all the cells were washed twice with ice cold PBS, harvested and counted under a phase-contrast microscope. Values are expressed as averages \pm S.D. (n=3).

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For [^3H]-thymidine incorporation assays, BR293 cells were seeded into 96 well plates (2×10^4 /well) in DME supplemented with 10% FCS and grown over night. The medium was then replaced with serum-free medium. Twenty-four hours later, the designated medium was added and after 18h, 0.5 μCi of (methyl)-[^3H]-thymidine (20 Ci mmol $^{-1}$, Amersham Corp.) was added to each well for an additional 18h. The cells were then harvested with trypsin/EDTA and transferred to a cellulose-coated plate and washed. Incorporated radioactivity was measured with a microplate scintillation counter. Values are expressed as averages \pm SD (n=4).

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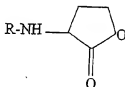
CLAIMS

1. An autocrine/paracrine signalling pathway which activates STAT wherein the pathway requires JAK activity and does not require ErbB1 activity and is not induced by EGF.
- 5 2. A process wherein STAT dimers accumulate in the cytoplasm wherein the process does not require ErbB1 activity or JAK activity.
3. The modulation of any one of the pathways or processes claimed in Claim 1 or Claim 2 to alter the amount of activated STAT.
- 10 4. The use of a compound selected from:-JAK, ErbB1, EGF, ErbB1inhibitors, EGF inhibitors, STAT inhibitors, interleukin- 13 (IL-13), IL-13E13K (IL-13 in which the Glu at position 13 is substituted by a Lys residue), sulphur methoxymethyl,ubiquitin E3 ligase, serine phosphatase, tyrosine phosphatase, SOCs, Pias proteins (protein inhibitors of activated
- 15 STAT), STAT1 inhibitors, STAT2 inhibitors, STAT3 inhibitors, STAT4 inhibitors, STAT5A inhibitors, STAT5Binhibitors, STAT6 inhibitors, JAK inhibitors, AG 490, α -amanitin, transcription inhibitors, quorum sensing molecules, N-acyl homoserine lactones, N-(3-oxododecanoyl)-L-homoserine lactone, oxygen radical scavengers, N-acetyl Cysteine (NAC), diphenylene iodonium chloride (DPI), inhibitors of COX1, inhibitors of COX2, aspirin,
- 20 ketorolac, indomethacin, or panCOX inhibitors to modulate STAT activity for the treatment of cancer, breast cancer, multiple myeloma, head and neck cancers, leukaemia, HTLV-1-dependent leukemia, large granular lymphocyte (LGL) leukaemia, erythroleukemia, acute lymphocytic leukemia (ALL), chronic lymphocytic leukaemia (CLL), acute myelogenous leukemia (AML), chronic myelogenous leukaemia (CML), megakaryoticleukaemia, lung
- 25 cancer, renal cell carcinoma, prostrate carcinoma, melanoma, ovarian carcinoma, pancreatic adenocarcinoma, lymphoma, EBV-related lymphoma, Burkitt's lymphoma mycosis fungoides lymphoma, HSV saimiri-dependent (T cell) lymphoma, cutaneous T cell lymphoma, obesity, lipid metabolism disorders, immune disease, immune deficiency or immune disorders.

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5. The use of a compound of the formula I



in which R is an acyl group of the formula II



wherein one of R^1 and R^2 is H and the other is selected from OR^4 , SR^4 and NHR^4 wherein R^4 is H or 1-6C alkyl, or R^1 and R^2 together with the carbon atom to which they are joined form a keto group and R^3 is a straight or branched chain saturated or unsaturated aliphatic hydrocarbonyl group containing from 8 to 11 carbon atoms and is optionally substituted by one or more substituent groups selected from halo, 1-6C alkoxy, carboxy, 1-6C alkoxycarbonyl, carbamoyl optionally mono- or disubstituted at the N atom by 1-6C alkyl and NR^5R^6 wherein each of the R^5 and R^6 is selected from H and 1-6C alkyl or R^5 and R^6 together with the N atom form a morpholino or piperazino group or any enantiomer thereof with the proviso that R is not a 3-oxododecanoyl group to modulate STAT activity for the treatment of cancer, breast cancer, multiple myeloma, head and neck cancers, leukaemia, HTLV-1-dependent leukemia, large granular lymphocyte (LGL) leukaemia, erythroleukemia, acute lymphocytic leukemia (ALL), chronic lymphocytic leukaemia (CLL), acute myelogenous leukemia (AML), chronic myelogenous leukaemia (CML), megakaryotic leukaemia, lung cancer, renal cell carcinoma, prostate carcinoma, melanoma, ovarian carcinoma, pancreatic adenocarcinoma, lymphoma, EBV-related lymphoma, Burkitt's lymphoma mycosis

fungoides lymphoma, HSV saimiri-dependent (T cell) lymphoma, cutaneous T cell lymphoma, obesity, lipid metabolism disorders, immune disease, immune deficiency or immune disorders.

6. The use claimed in claim 5 wherein the R group is selected from

5



10 and



15

wherein R³ is as defined in claim 5.

7. The use claimed in either claim 5 or claim 6 wherein the group R³ is an 8-11C straight or branched chain alkyl group optionally substituted by a substituent selected from

20 bromo, carboxy and methoxycarbonyl.

8. The use claimed in any one of claims 5 to 7 wherein the R³ group is such that the group R in formula I is selected from;

25

3-oxoundecanoyl;

11-bromo-3-oxoundecanoyl;

10-methyl-3-oxoundecanoyl;

6-methyl-3-oxoundecanoyl;

3-hydroxydodecanoyl;

30

12-bromo-3-oxododecanoyl;

- 3-oxotridecanoyl;
13-bromo-3-oxotridecanoyl;
3-hydroxytetradecanoyl; 3-oxotetradecanoyl;
14-bromo-3-oxotetradecanoyl; and
5 13-methoxycarbonyl-3-oxotridecanoyl.

9. The use claimed in any one of claims 5 to 8 wherein the R^3 is an 8-11 straight or branched chain alkenyl group optionally substituted by a substituent selected from bromo, carboxy and methoxycarbonyl.

10

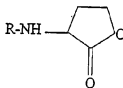
10. The use claimed in any one of claims 5 to 9 wherein the R^3 group is such that the group R in formula I is selected from;

- 3-oxo-12-tridecenoyl;
3-oxo-7-tridecenoyl; 3-hydroxy-7-tetradecenoyl;
15 3-oxo-9-tetradecenoyl;
3-hydroxy-9-tetradecenoyl;
3-oxo-10-tetradecenoyl;
3-hydroxy-10-tetradecenoyl; 3-oxo-11-tetradecenoyl;
3-hydroxy-11-tetradecenoyl;
20 3-oxo-13-tetradecenoyl; and
3-hydroxy-13-tetradecenoyl.

11. The use of JAK, ErbB1, EGF, ErbB1 inhibitors, EGF inhibitors, STAT inhibitors, interleukin-13 (IL-13), IL-13E13K (IL-13 in which the Glu at position 13 is substituted by a
25 Lys residue), sulphur methoxymethyl, ubiquitin E3 ligase, serine phosphatase, tyrosine phosphatase, SOCs, Pias proteins (protein inhibitors of activated STAT), STAT1 inhibitors, STAT2 inhibitors, STAT3 inhibitors, STAT4 inhibitors, STAT5A inhibitors, STAT5B inhibitors, STAT6 inhibitors, JAK inhibitors, AG 490, α -amanitin, transcription inhibitors, quorum sensing molecules, N-acyl homoserine lactones, N-(3-oxododecanoyl)-L-homoserine
30 lactone, oxygen radical scavengers, N-acetyl Cysteine (NAC), diphenyltetrabutylammonium chloride

(DPI), inhibitors of COX1, inhibitors of COX2, aspirin, ketorolac, indomethacin, or panCOX inhibitors for the preparation of a medicament for the treatment of cancer, breast cancer, multiple myeloma, head and neck cancers, leukaemia, HTLV-1-dependent leukemia, large granular lymphocyte (LGL) leukaemia, erythroleukemia, acute lymphocytic leukaemia(ALL), chronic lymphocytic leukaemia (CLL), acute myelogenous leukemia (AML), chronic myelogenous leukaemia (CML), megakaryotic leukaemia, lung cancer, renal cell carcinoma, prostate carcinoma, melanoma, ovarian carcinoma, pancreatic adenocarcinoma, lymphoma, EBV-related lymphoma, Burkitt's lymphoma mycosis fungoides lymphoma, HSV saimiri-dependent (T cell)lymphoma, cutaneous T cell lymphoma, obesity, lipid metabolism disorders, immune disease, immune deficiency or immune disorders.

12. The use of a compound of the formula I



in which R is an acyl group of the formula II



wherein one of R^1 and R^2 is H and the other is selected from OR^4 , SR^4 and NHR^4
 wherein R^4 is H or 1-6C alkyl, or R^1 and R^2 together with the carbon atom to which they are joined form a keto group and R^3 is a straight or branched chain saturated or unsaturated aliphatic hydrocarbyl group containing from 8 to 11 carbon atoms and is optionally substituted by one or more substituent groups selected from halo, 1-6C alkoxy, carboxy, 1-6C alkoxy carbonyl, carbamoyl optionally mono- or disubstituted at the N atom by 1-6C alkyl and NR^5R^6 wherein each of the R^5 and R^6 is selected from H and 1-6C alkyl or R^5 and R^6

together with the N atom form a morpholino or piperazino group or any enantiomer thereof with the proviso that R is not a 3-oxododecanoyl group for the preparation of a medicament for the treatment of treatment of cancer, breast cancer, multiple myeloma, head and neck cancers, leukaemia, HTLV-1-dependent leukemia, large granular, lymphocyte
 5 (LGL) leukaemia (ALL), chronic lymphocytic leukaemia (CLL), acute myelogenous leukemia (AML), chronic myelogenous leukaemia (CML), megakaryotic leukaemia, lung cancer, renal cell carcinoma, prostate carcinoma, melanoma, ovarian carcinoma, pancreatic adenocarcinoma, lymphoma, EBV-related lymphoma, Burkitt's lymphoma mycosis fungoides lymphoma, HSV saimiri-dependent (T cell) lymphoma, cutaneous T cell
 10 lymphoma, obesity, lipid metabolism disorders, immune disease, immunodeficiency or immune disorders.

13. The use claimed in claim 12 wherein the R group is selected from

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and

20



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wherein R³ is as defined in claim 12.

14. The use claimed in either claim 12 or claim 13 wherein the group R³ is an 8-11C straight or branched chain alkyl group optionally substituted by a substituent selected from
 30 bromo, carboxy and methoxycarbonyl.

15. The use claimed in any one of claims 12 to 14 wherein the R³ group is such that the group R in formula I is selected from;

- 3-oxoundecanoyl;
- 11-bromo-3-oxoundecanoyl;
- 5 10-methyl-3-oxoundecanoyl;
- 6-methyl-3-oxoundecanoyl;
- 3-hydroxydodecanoyl;
- 12-bromo-3-oxododecanoyl;
- 3-oxotridecanoyl;
- 10 13-bromo-3-oxotridecanoyl;
- 3-hydroxytetradecanoyl;
- 3-oxotetradecanoyl;
- 14-bromo-3-oxotetradecanoyl; and
- 13-methoxycarbonyl-3-oxotridecanoyl.

15

16. The use claimed in any one of claims 12 to 15 wherein the R³ is an 8-11 straight or branched chain alkenyl group optionally substituted by a substituent selected from bromo, carboxy and methoxycarbonyl.

20 17. The use claimed in any one of claims 12 to 16 wherein the R³ group is such that the group R in formula I is selected from;

- 3-oxo-12-tridecenoyl;
- 3-oxo-7-tridecenoyl;
- 25 3-hydroxy-7-tetradecenoyl;
- 3-oxo-9-tetradecenoyl;
- 3-hydroxy-9-tetradecenoyl;
- 3-oxo-10-tetradecenoyl;
- 3-hydroxy-10-tetradecenoyl;
- 30 3-oxo-11-tetradecenoyl;

3-hydroxy-11-tetradecenyl;
3-oxo-13-tetradecenyl; and
3-hydroxy-13-tetradecenyl.

Figure 1

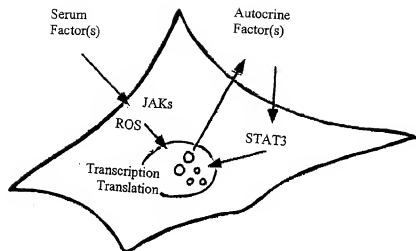


Figure 2

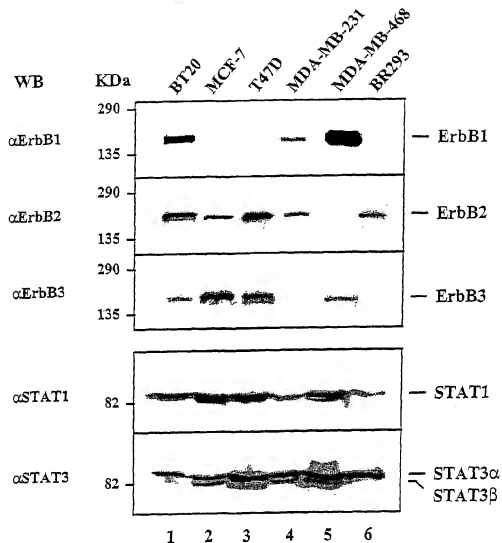


Figure 3

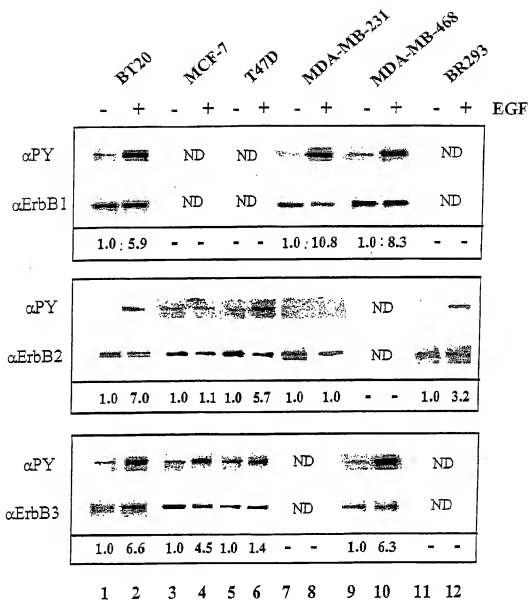


Figure 4

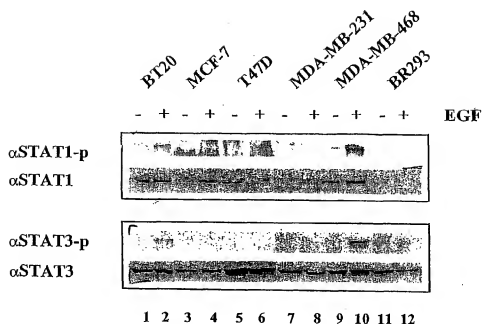


Figure 5

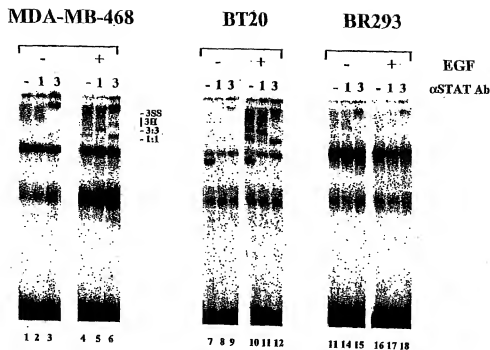
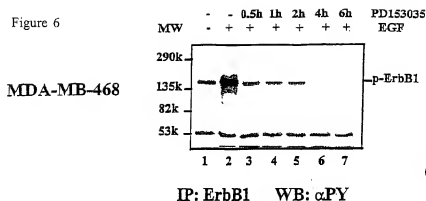
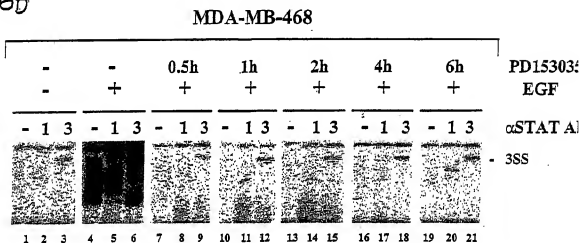


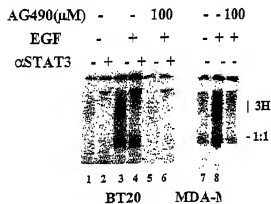
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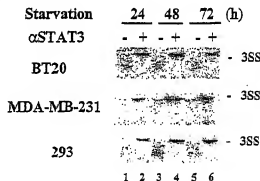
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6c



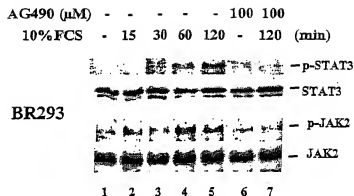
6d



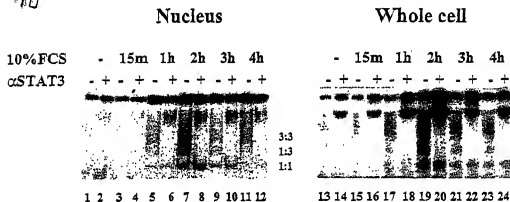
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Figure 7

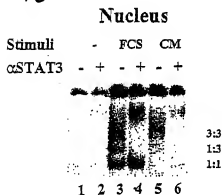
7a



7b



7c



7d

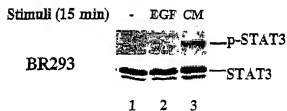
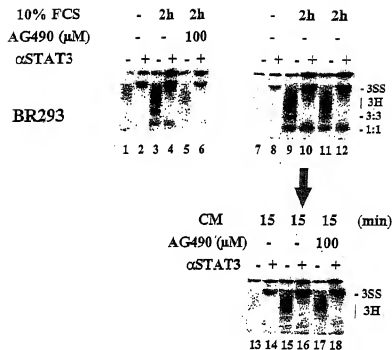


Figure 8

8a



8b

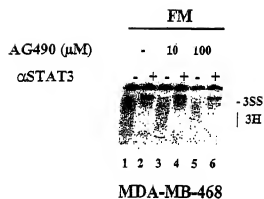


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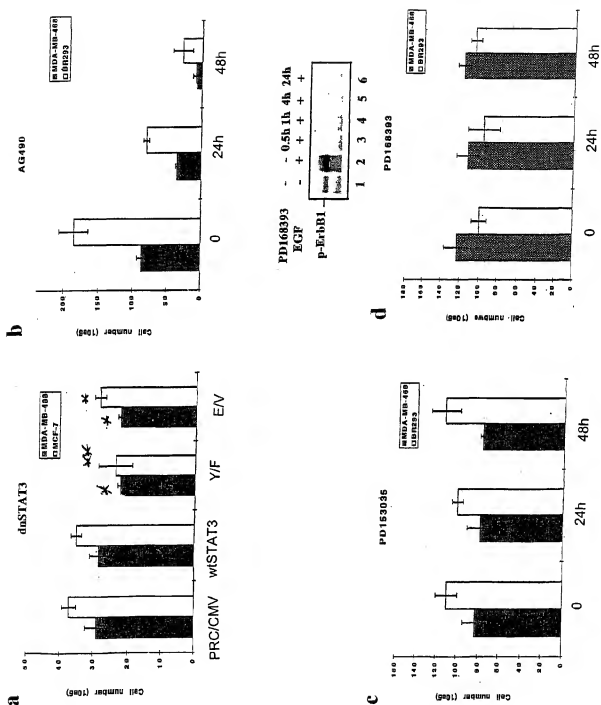
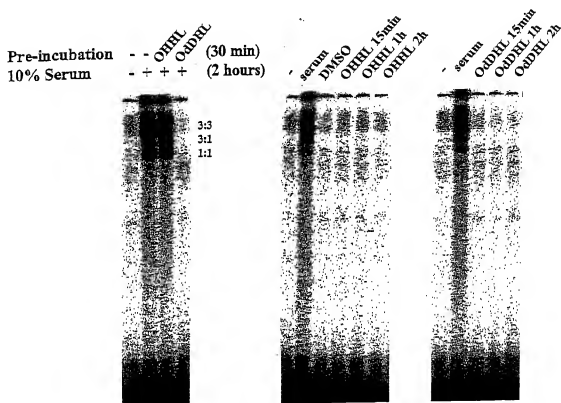


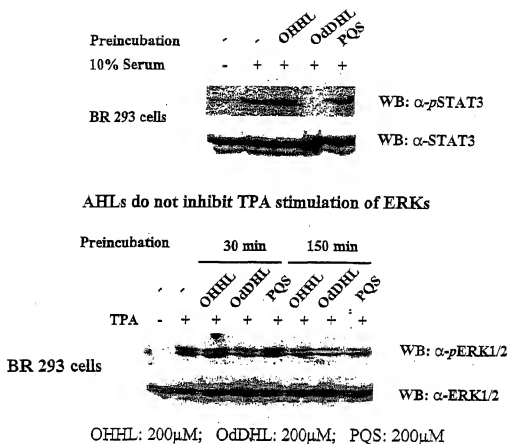
Figure 10

OdDHL blocks stimulation of STAT1 and 3 by serum



BR 293 cells
SIE/M67 probe
OHHL and OdDHL: 200 μ M

Figure 11



ROS scavengers and AG490 inhibit serum stimulation of STATs

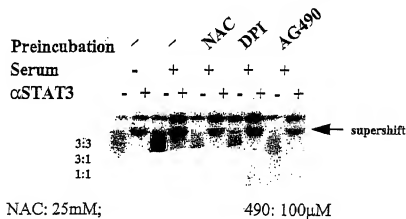


Figure 12

Dose response of OdDHL effect on STAT3 DNA binding (left).
 OdDHL inhibits STAT3 activation by H_2O_2 (middle) but not
 activation by conditioned medium (CM).

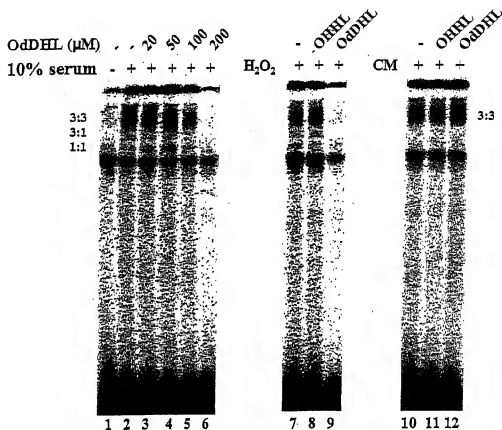
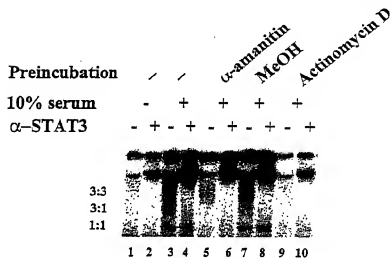


Figure 13

Serum stimulation of STAT1 and 3 is blocked by the transcription inhibitors α -amanitin and actinomycin D



OdDHL inhibits generation of serum stimulated autocrine factors in BR 293 cells

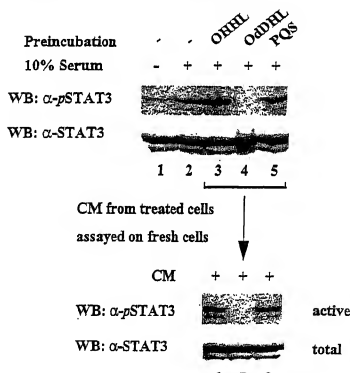


Figure 14

**OdDHL blocks proliferation (48h) and
induces apoptosis (24h) in BR293 cells**



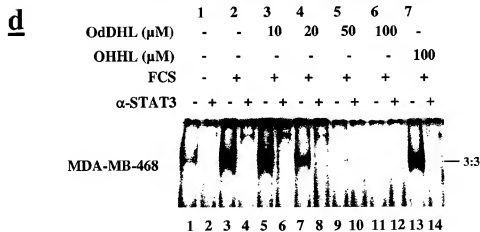
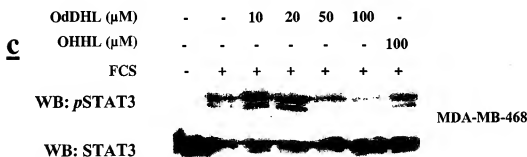
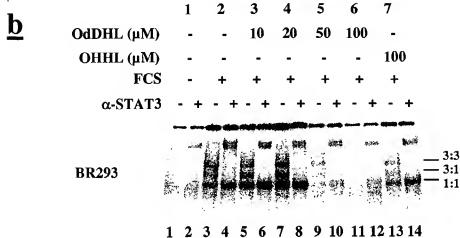
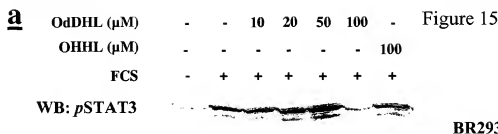
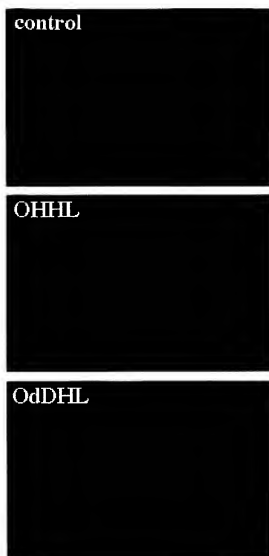
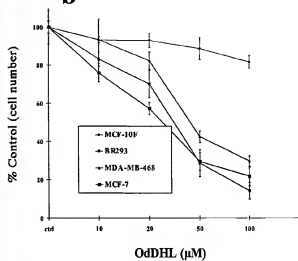
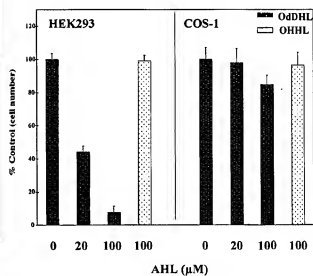
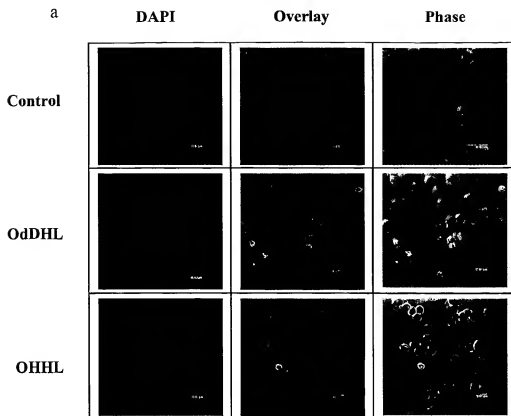


Figure 16

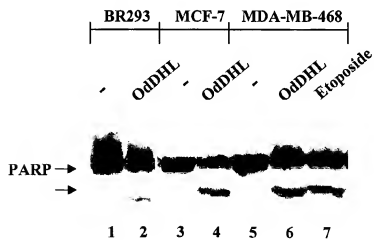
a**b****c**

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Figure 17



b



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Figure 18

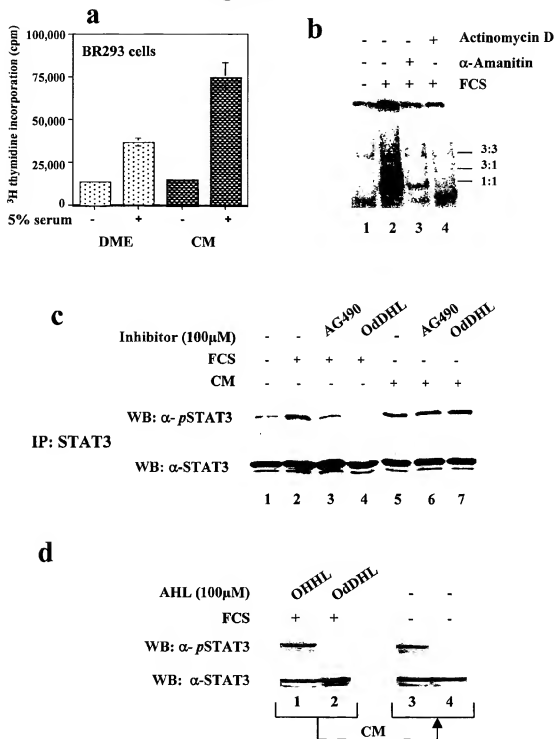


Figure 19

